## **AMENDMENTS TO THE SPECIFICATIONS**

Please amend Table 4 on page 75 as follows:

**TABLE 4: PCR Primers** 

Reference SNP ID	Forward PCR primer	SEQ ID NO.	Reverse PCR primer	SEQ ID NO.
rs1671152	ACGTTGGATGAGGGCTGTGCAGAGGCCGCTT	<u>18</u>	ACGTTGGATGTGAACATCCTGTCGGCCTCC	<u>19</u>
rs1050348	CAGCTGGATGACTACAATGC	<u>20</u>	TGTTCATGTCTTCGGCATCC	<u>21</u>
rs454422	CAGCTTTTGAGGCACTTTCC	22 AGCACCTTGCATACCCATAG		<u>23</u>
rs763471	TAACTCCTGTGTGGCTTTCT	<u>24</u>	GTGAAGAGCTCTGAAATGCC	<u>25</u>
rs2046778	CATGAAGCCTTATGCTTGAG	<u>26</u>	GTTCCCTTCCCCCATAAAAC	<u>27</u>

Please amend Table 5 on page 76 as follows:

**TABLE 5: Extend Primers** 

Reference SNP ID	Extend Probe	SEQ ID NO.	Term Mix
rs1671152	CTCCATCCTGACCCCCGT	<u>28</u>	ACT
rs1050348	CACTTGACCAGGCCCTTAAC	<u>29</u>	ACG
rs454422	GATCCTTCTCACTTACTGTTC	<u>30</u>	ACT
rs763471	CTCCAAGCAGTAAAGATGTTC	<u>31</u>	CGT
rs2046778	CTGTCATGATTGACAGGTCC	<u>32</u>	ACT

Please amend Table 8 on pages 82-84 as follows:

**TABLE 8** 

dbSNP rs#	Forward PCR primer	SEQ ID	Reverse PCR primer	SEQ ID
10666	ACGTTGGATGAGATGGCCCCTCTCCCCT	<u>33</u>	ACGTTGGATGAGTGGACTGGCCTGCAGGT	34
172006	ACGTTGGATGGGTTGAGGAGTATTCCATTG	<u>35</u>	ACGTTGGATGTGGGTGACAGCAAGACTCCA	<u>36</u>
269909	ACGTTGGATGAGTTTCTTGTCTCCTGGGTG	<u>37</u>	ACGTTGGATGCACAACAATGAAAGGACAAGC	<u>38</u>
269910	ACGTTGGATGGGCTGCTCAGGTTCTAAAAG	<u>39</u>	ACGTTGGATGCCCCAGTTCCTTATCTGATC	<u>40</u>
269911	ACGTTGGATGGGTGACAAAGTGAGACTCCG	41	ACGTTGGATGTGACTAGCTGGGATTATGGG	42
269912	ACGTTGGATGAGGAGAGCCTGCAGGTTGAA	<u>43</u>	ACGTTGGATGTCCCTTGATTGTCATCACAG	44
269913	ACGTTGGATGAGATGCACCGAATGGATCTG	<u>45</u>	ACGTTGGATGTCCTAGGACACAGTGTGGAC	<u>46</u>
269915	ACGTTGGATGAAGCTGAGATTGTGCTGCTG	<u>47</u>	ACGTTGGATGACTACTCTACTTTCTACCCC	<u>48</u>
269916	ACGTTGGATGAACTCCTGACCACGTGATCC	<u>49</u>	ACGTTGGATGAAAGTGTAGCTGGGCATGGC	<u>50</u>
703464	ACGTTGGATGAAAAGTAGCTGAGCGTGGTG	<u>51</u>	ACGTTGGATGCCCAGGTTCAAACAGTTCTC	<u>52</u>
703465	ACGTTGGATGGCTGTGATGACAATCAAGGG	<u>53</u>	ACGTTGGATGAGCAGGAACTGGCATTTGAG	<u>54</u>
703467	ACGTTGGATGAAACTTACGAAGGTCTGGGC	<u>55</u>	ACGTTGGATGGGGTTTCAGGAACATTCACC	<u>56</u>

dbSNP rs#	Forward PCR primer	SEQ ID NO.	Reverse PCR primer	SEQ ID NO.
703468	ACGTTGGATGGAAAGGAACAAGTGATCCAG	<u>57</u>	ACGTTGGATGCTTCTGAAAAAGGAGAAGGG	<u>58</u>
754235	ACGTTGGATGGTGGAACAAACAGTTGGAGC	<u>59</u>	ACGTTGGATGGAGTGGAGATCATGCCATTG	<u>60</u>
775821	ACGTTGGATGTTTTCTCATCCGCCAGCAGG	<u>61</u>	ACGTTGGATGACAGAGCACAGGTCCCTTTC	<u>62</u>
775822	ACGTTGGATGAAAAGTAGAGCATGTGGACC	63	ACGTTGGATGGGGATGAACAGACAATCCTC	<u>64</u>
775894	ACGTTGGATGAGGCAGGAGGACTGATTGAG	<u>65</u>	ACGTTGGATGTTGTAGAGACAGGGTCTTGC	<u>66</u>
775900	ACGTTGGATGTCTGCAACGTTCGTTCTTCC	67	ACGTTGGATGTTCTTAATGTGCCCACTGTC	<u>68</u>
775903	ACGTTGGATGTGTGCCCGGCCCTTTTTTTC	69	ACGTTGGATGGGACTCCAGTCTGAGTGACA	<u>70</u>
776251	ACGTTGGATGAATCCTAGCTACTCAGGAGG	<u>71</u>	ACGTTGGATGATGGTATGATCTAGGCTCCC	<u>72</u>
892088	ACGTTGGATGATTGGTTACACTGGGACTGC	<u>73</u>	ACGTTGGATGAAGCGGTGATTCTCAGCCTC	<u>74</u>
892089	ACGTTGGATGCCCTACAAGAATCCCGAGAG	<u>75</u>	ACGTTGGATGAAGCTGTAGCATCGGTAGGTT	<u>76</u>
892090	ACGTTGGATGAAGCTGTAGCATCGGTAGGTT	77	ACGTTGGATGCCCTACAAGAATCCCGAGAG	<u>78</u>
892091	ACGTTGGATGAAGATGGAGGACCACAGGTG	<u>79</u>	ACGTTGGATGACCGGAATTACTGAGAGGTC	80
1036231	ACGTTGGATGAACAGTACTAAGGGCAGATG	<u>81</u>	ACGTTGGATGGTCCAGGGTGTTTACTGTTC	<u>82</u>
1036232	ACGTTGGATGCCAAAGAAACTCCCAGAATC	<u>83</u>	ACGTTGGATGGATCGTGCCATTGCACTTTG	<u>84</u>
1043673	ACGTTGGATGTGAAGTCATGAGAAGAAGGC	<u>85</u>	ACGTTGGATGGCTGCTGGAAGAAATAGAAG	<u>86</u>
1043678	ACGTTGGATGTTCATGATCTGAATCCCCCC	<u>87</u>	ACGTTGGATGGAACTCATTATCCCTGAGGG	<u>88</u>
1043680	ACGTTGGATGTCTAGCCCAGCAATGAACTC	<u>89</u>	ACGTTGGATGTTCCAGGTGTTGGTGAACTG	<u>90</u>
1043684	ACGTTGGATGCTCAATATACCCGTGATACAG	<u>91</u>	ACGTTGGATGTTTAGCCATGATTCTGCCTC	<u>92</u>
1054796	ACGTTGGATGAGTGTCACCCTGATTTCCAG	<u>93</u>	ACGTTGGATGCACCTTGGGAAATACGTTGC	<u>94</u>
1059211	ACGTTGGATGCTCAGCTCCTTGGTGAAGAG	<u>95</u>	ACGTTGGATGGGCAGACGAGGAAGTATAAC	<u>96</u>
1064675	ACGTTGGATGGAGTTCCCTCAGTTTTTATTG	<u>97</u>	ACGTTGGATGCCTACTACATTCCTTTTTGC	<u>98</u>
1560714	ACGTTGGATGATCTGCTGACCTCGTGATCC	99	ACGTTGGATGAAAAGACAGTCTCAGGTGGG	<u>100</u>
1613662	ACGTTGGATGATGGACCCTGCAGAACCTAC	<u>101</u>	ACGTTGGATGTCTGATTTCCCAGGAACCTC	<u>102</u>
1625609	ACGTTGGATGTCAAGCGATTCTCCTGCCTC	<u>103</u>	ACGTTGGATGAAAAAATGAGCTGGGCGTGG	104
1625689	ACGTTGGATGGTAATCCCAGCTACTTGGAG	<u>105</u>	ACGTTGGATGATCTTGGCTCACTGCAGCCT	<u>106</u>
1626971	ACGTTGGATGTATTAAATGCACCTGGCACC	107	ACGTTGGATGCAAAGTGCTGGGATTACAGG	<u>108</u>
1654406	ACGTTGGATGCTTCTATTTCTTCCAGCAGC	<u>109</u>	ACGTTGGATGTTTCTTCCCCCATTGTACCC	<u>110</u>
1654409	ACGTTGGATGGTGAAACCTTGTCTCATATAC	111	ACGTTGGATGTGAGAGTAGGCATGTGGTAC	112
1654410	ACGTTGGATGACTGTGCCTAGGCTATACTG	113	ACGTTGGATGGGAAAATCATACTGAGATGC	114
1654411	ACGTTGGATGTGGAACTTCTTGGTGCCATC	<u>115</u>	ACGTTGGATGCCTGTAATCCCGGCACTTTG	<u>116</u>
1654412	ACGTTGGATGATTAGCCAGGTGTGGTGGTG	117	ACGTTGGATGTCAAGCCATTCTCCCACCTC	<u>118</u>
1654413	ACGTTGGATGAGGGCTGTGCAGAGGCCGCTT	<u>119</u>	ACGTTGGATGTCCATCCTGACCCCCGTTTG	120
1654415	ACGTTGGATGCCAAGAAAGTCCTTGGTGTG	121	ACGTTGGATGCTTTGAAATGGCCCCATCAC	122
1654416	ACGTTGGATGTCTGCTGAGCATGAAATGCC	<u>123</u>	ACGTTGGATGCTGAACTGACCGTCTCATTC	<u>124</u>
1654419	ACGTTGGATGTATCATACGCTAGGCTGGAG	<u>125</u>	ACGTTGGATGATGTTTCTCCTGCCTTGGTG	<u>126</u>
1654420	ACGTTGGATGCCAACCAACCAACAACCTG	<u>127</u>	ACGTTGGATGTGGAAGTTTGAGAACCGCTG	<u>128</u>
1654421	ACGTTGGATGAGGACACAGGAATCCAGAAG	<u>129</u>	ACGTTGGATGGCACATTCTGGGCTATTAAC	<u>130</u>
1654424	ACGTTGGATGTAGGTGGGAAGGAAGTGGGA	<u>131</u>	ACGTTGGATGCCACTTCTTTCCCACCTATG	<u>132</u>
1654425	ACGTTGGATGTACCTGTGACCACAAGCTCC	<u>133</u>	ACGTTGGATGTGCTACAGCTTCTCCAGCAG	<u>134</u>
1654438	ACGTTGGATGAATCAACTAGGCATGGTGGC	<u>135</u>	ACGTTGGATGCCAGGTTCAAGCGATTCTCC	<u>136</u>
1654439	ACGTTGGATGCCCCATATACATGTGCGATG	<u>137</u>	ACGTTGGATGAATGGGGTGTTTCTGGAGCA	<u>138</u>
1654441	ACGTTGGATGAGTAGCTGGGATTACAGGCG	<u>139</u>	ACGTTGGATGGGAGTTCAAGATAAGCCTGG	<u>140</u>
1654442	ACGTTGGATGAGGAGAATGGTGTGAAGCTG	141	ACGTTGGATGAATCTTGCTCTGTCACCCAG	<u>142</u>
1654444	ACGTTGGATGGGATGGTCCCAGTTTTACAT	143	ACGTTGGATGCCAGGAGAATCACTTTTATGG	
1654446	ACGTTGGATGAAAAGGAAGGGCATTCTGGC	<u>145</u>	ACGTTGGATGTTTGGCCTCCCAAAGTACTG	<u>146</u>
1654447	ACGTTGGATGATCCCTGGGAAGACGGTCAT	147	ACGTTGGATGTTACCTCTCCTGGCCAGTTC	<u>148</u>

dbSNP		SEQ ID		SEQ ID
rs#	PCR primer	<u>NO.</u>	PCR primer	<u>NO.</u>
1654448	ACGTTGGATGTGCTCACTGCATGAGATTCC	149	ACGTTGGATGAACTTTGGCCTCCCAAAGTG	<u>150</u>
1654449	ACGTTGGATGAGTCCAGCCTGGCAAACATG	<u>151</u>	ACGTTGGATGCAGTCTAATCTCTCTTTTCCC	<u>152</u>
1654451	ACGTTGGATGTTTAAAATGCCCGCTGCACG	<u>153</u>	ACGTTGGATGAGGAGGATGCACTTATGTGG	<u>154</u>
1654452	ACGTTGGATGCTGTACGCATTACCACAGAC	<u>155</u>	ACGTTGGATGGTTTTTGGACTCTTGACCTGC	<u>156</u>
1654459	ACGTTGGATGCAGGAGCTTGGGTACCCAC	157	ACGTTGGATGCCCTCATCTGGAAATGTGTG	<u>158</u>
1654485	ACGTTGGATGTTGTACCACTGCACTCTAGC	<u>159</u>	ACGTTGGATGCCTGACTCTACAGTTCTTGC	<u>160</u>
1654491	ACGTTGGATGCAGACGTCCGTGCTTCACC	<u>161</u>	ACGTTGGATGTCCAGGAACAGACGGAGGTC	<u>162</u>
1654495	ACGTTGGATGATGACCATTGCTCGTCTGTG	<u>163</u>	ACGTTGGATGGCTTTCTGCAGAGGTTGTCG	<u>164</u>
1654496	ACGTTGGATGAATCACAAATGGCAACACGG	<u>165</u>	ACGTTGGATGTTTGGATGCTGGCACTTGTG	<u>166</u>
1654497	ACGTTGGATGACCCCATGCTGTGTTTTCTC	<u>167</u>	ACGTTGGATGCAGAAGACTACCTGATTTGC	<u>168</u>
1654498	ACGTTGGATGCTTCCCACACCCACTATATC	<u>169</u>	ACGTTGGATGGTTAGTGAGTCGGTGACATC	<u>170</u>
1654499	ACGTTGGATGCACTACCTCTCTAGCAACTG	171	ACGTTGGATGACCTCAGATGATCTGCCCAC	<u>172</u>
1654503	ACGTTGGATGTCCTTGGCTTGTGGCCCTTC	<u>173</u>	ACGTTGGATGAGCCAGGGCAACGTTTGAAG	174
1654504	ACGTTGGATGCCACCCCATGATTCCATTTC	<u>175</u>	ACGTTGGATGTGCTGTGATGCACCTTTGAC	<u>176</u>
1654505	ACGTTGGATGCCCTGTCTCTCTAAAACCAC	177	ACGTTGGATGATTCAAGCAGTTCTCGTGCC	<u>178</u>
1671133	ACGTTGGATGGTGGTCTCAACTTGGCTATC	<u>179</u>	ACGTTGGATGCCAGATAGGATTCCAGGTTC	<u>180</u>
1671140	ACGTTGGATGAGTCTGACAAGAGAGTCAGC	<u>181</u>	ACGTTGGATGTCCTTTACCTACCCACATCC	<u>182</u>
1671148	ACGTTGGATGGCCATCCTTCTGTCTTTTCC	<u>183</u>	ACGTTGGATGAGTGGCTCATGCCTGTAATC	184
1671149	ACGTTGGATGCTTTTCCCAAGTGACTCACC	185	ACGTTGGATGAAAAGAATGGCTGGCCACAG	<u>186</u>
1671150	ACGTTGGATGGTGCTATGATCAAATCAGGG	187	ACGTTGGATGACACCACTGCACTCTAGCTC	188
1671151	ACGTTGGATGGGAAAACCAGACAAGAGCAC	189	ACGTTGGATGTGACTCTGTTCCATCCTCTG	<u>190</u>
	ACGTTGGATGAGGGCTGTGCAGAGGCCGCTT	191	ACGTTGGATGTGAACATCCTGTCGGCCTCC	<u>192</u>
1671153	ACGTTGGATGCCTACTCCGAACACACACAC	<u>193</u>	ACGTTGGATGATTATAGGCATGAGGCACCG	<u>194</u>
1671169	ACGTTGGATGTCCTGTTGCTGGACACTATC	<u>195</u>	ACGTTGGATGTCACACCTTCCGAGGATTAG	<u>196</u>
1671170	ACGTTGGATGAGGTGACAGTGCTGTACCTG	<u>197</u>	ACGTTGGATGACAAAGAACAGTGAGAGGGC	<u>198</u>
1671171	ACGTTGGATGAAGCAAGATACCGTCTCAGA	<u>199</u>	ACGTTGGATGCCGGGAAATGGAATAATTCC	<u>200</u>
1671176	ACGTTGGATGTGGAGCCACTTATGGAGAAC	<u>201</u>	ACGTTGGATGACCCCAACTGAAACACAGAC	202
1671178	ACGTTGGATGTAATCCCAGCACTTTGGGAG	<u>203</u>	ACGTTGGATGCATGTTTGCCAGGCTGGACT	<u>204</u>
1671182	ACGTTGGATGATAGGGCGGCTTTTCTCCTG	<u>205</u>	ACGTTGGATGCCTGGGAACTGAATGTCTCG	<u>206</u>
1671187	ACGTTGGATGAGTGCTCAGCAACGATTACG	207	ACGTTGGATGGAGGGCTGCAGGTTGAGAAA	<u>208</u>
1671188	ACGTTGGATGGGAACCGCAGATGGACAATG	<u>209</u>	ACGTTGGATGAGATCACAGAGTGAGGAGAG	210
	ACGTTGGATGTCGGACGCACACAGACTGTAG	<u>211</u>	ACGTTGGATGGGAAAGCGTATCTGCAGAGG	<u>212</u>
1671192	ACGTTGGATGTGGTAAGAGACGGACAGTTC	<u>213</u>	ACGTTGGATGTCAGCAGAAAGGAGTGTGAG	
1671196	ACGTTGGATGTTGCTAGGCAACAGGCACTC	<u>215</u>	ACGTTGGATGTCTGTATCTGAGCCTCACTG	<u>216</u>
1671198	ACGTTGGATGATGAAACTAAGGCACATGGC	217	ACGTTGGATGCTTATAATCTACCCTCTTAGC	<u>218</u>
1671199	ACGTTGGATGGCTGAAATTTGCTAAGAGGG	<u>219</u>	ACGTTGGATGGACAGTTACTACTAGCAAGC	<u>220</u>
1671214	ACGTTGGATGAGGCGGAGAATGATCCGGTG	<u>221</u>	ACGTTGGATGACGCCATCATTCGTGCATCC	<u>223</u>
1671215	ACGTTGGATGTTCTCCAAAGCACCCAAGTG	<u>224</u>	ACGTTGGATGATGCTGGGCTTGCTTTTTCC	<u>225</u>
1671216	ACGTTGGATGTGCTTGGGAGCAAGTTACAG	<u>226</u>	ACGTTGGATGTTCCCCCTCCTGGTATTTAC	<u>227</u>
1671217	ACGTTGGATGTTGTCTCCATTCCTCCCTGG	228	ACGTTGGATGTCTTGTCTTGCCCTCTCGCT	<u>229</u>
1671218	ACGTTGGATGTGAGTCTGGTAGGCAACTTC	<u>230</u>	ACGTTGGATGTAGAAGCCAGTCGCTACATC	<u>231</u>
1671219	ACGTTGGATGTGATCTCGGCTCACTGCAAG	<u>232</u>	ACGTTGGATGAAATTAGCTGGGCATGGTGG	<u>233</u>
1671221	ACGTTGGATGTGGTGAAACCCCATTTCTAC	<u>234</u>	ACGTTGGATGGGTTCAAGGGATTCTCCTGC	<u>235</u>
1671223	ACGTTGGATGTCAAGTGATTCTCCTGCCTC	236	ACGTTGGATGCCCACCTCTACTGAAAATAC	237
1671224	ACGTTGGATGTGAGTCTCACTCTTGTTGCC	<u>238</u>	ACGTTGGATGCAGGAGAATCACTTGAACCC	<u>239</u>
1671225	ACGTTGGATGTATAGGCGTGAGCCACTATG	<u>240</u>	ACGTTGGATGCTATTGGAAGCTACATGCTC	241

dbSNP rs#	Forward PCR primer	SEQ ID NO.	Reverse PCR primer	SEQ ID NO.
1671226	ACGTTGGATGTATTGGCCAGACTGGACTTC	241	ACGTTGGATGAGTTACTCAGGAGGCTAAGG	242
1671227	ACGTTGGATGGGTTTCTGTTCAGAGATTCG	<u>243</u>	ACGTTGGATGTGCAGTGAGCCTAGATCATG	244
1671228	ACGTTGGATGTCAGCCTCCCAGGGATTAAG	<u>245</u>	ACGTTGGATGACATGGTGAAAACTCGTCTC	<u>246</u>
1869616	ACGTTGGATGTAATCCCAGCTACTCGGAAG	247	ACGTTGGATGACGGTGGCTCACTTCAACCT	<u>248</u>
2019599	ACGTTGGATGGTGCTGGGATTATAGGCATG	<u>249</u>	ACGTTGGATGTACTCCGAACACACACACAC	<u>250</u>
2116883	ACGTTGGATGATTACAGGCATGAGCCACTG	<u>251</u>	ACGTTGGATGCACGCGCAGTTCAATTTCTC	<u>252</u>
2124090	ACGTTGGATGTCTGACAAAGCTGGAAGCTG	<u>253</u>	ACGTTGGATGCTGATAAACAAGGCTGTGGG	<u>254</u>
2163833	ACGTTGGATGGATATTGGTGAGTATGCAGAG	<u>255</u>	ACGTTGGATGAACTGTTTTCCACAGCAGGG	<u>256</u>
2217659	ACGTTGGATGTTCCCCCCTTCTCCTTTTTC	<u>257</u>	ACGTTGGATGATGAGGTAACTTACCTAATG	<u>258</u>
2304167	ACGTTGGATGGTTTGGTTCCCAGAGACTTC	<u>259</u>	ACGTTGGATGAGGATGACTTACTCACCAGC	<u>260</u>
2304168	ACGTTGGATGTCAGCAGAAAGGAGTGTGAG	<u>261</u>	ACGTTGGATGTGGTAAGAGACGGACAGTTC	<u>262</u>
2365593	ACGTTGGATGTGACGCAGTAAGACTCCATC	<u>263</u>	ACGTTGGATGCAAAGTGCTGGGATTACAGG	<u>264</u>
2365721	ACGTTGGATGTTGTACAGCCTGCAAGCAAC	<u>265</u>	ACGTTGGATGAGATCGCGCCATTGCACTCA	<u>266</u>
2569513	ACGTTGGATGGTTGGCGTTTTTGTTTGCAC	<u>267</u>	ACGTTGGATGTCTCATAGTATTCTGCAGGG	<u>268</u>
2569514	ACGTTGGATGTCCCTGCAGAATACTATGAG	<u>269</u>	ACGTTGGATGAGAGTGTTGGGATTACAGGC	270
2886414	ACGTTGGATGGGTGTGCTTTACAAATGCTG	271	ACGTTGGATGAACTGAGATCACTCCACTGC	272
2886415	ACGTTGGATGTGACGCAGTAAGACTCCATC	<u>273</u>	ACGTTGGATGCAAAGTGCTGGGATTACAGG	274
3745912	ACGTTGGATGACGTCTTCTGAGGCACAGAG	<u>275</u>	ACGTTGGATGGCTGTTAGAGGCTGGCAGG	<u>276</u>
3786863	ACGTTGGATGTGACCAACAGAAGTCTCAGG	<u>277</u>	ACGTTGGATGTTGACCTCAGGTGATCCATC	<u>278</u>

Please amend Table 9 on pages 84-87 as follows:

TABLE 9

dbSNP rs#	Extend Primer	SEQ ID NO.	Term Mix
10666	TGCAGGTGAGCACTGCCC	<u>279</u>	ACG
172006	GCAAGACTCCATCTCAA	<u>280</u>	ACT
269909	AAGCATAGATCAGATAAGGAA	<u>281</u>	ACT
269910	ATCTATGCTTGTCCTTTCAT	<u>282</u>	ACT
269911	GCTCAGCTACTTTTTGTAT	<u>283</u>	CGT
269912	CAAGATGGTGTCTTCGGC	<u>284</u>	ACT
269913	ACAGTGTGGACCGATTTCC	<u>285</u>	ACT
269915	AGACAAGTCTCACTCTG	<u>286</u>	ACG
269916	GGCGGCTCACACCTGTAAT	<u>287</u>	ACG
703464	CTCCTGCCTCAGCCTCC	288	ACG
703465	TGGCATTTGAGACAGGA	<u>289</u>	ACT
703467	CATTCACCATGTCTGTGAG	<u>290</u>	ACG
703468	CTTCATAAAAGAAAGATGACA	<u>291</u>	ACG
754235	CATGCCATTGTACTCCAGCC	292	ACG
775821	CCTGCCAGCCTCTAACAGC	293	ACG
775822	TAGTGATGTCTGCTTCAG	<u>294</u>	ACT
775894	TTGCCCAGGCTGGCCTC	<u>295</u>	ACT
775900	GAATGCCAACCTCCCTTCC	<u>296</u>	ACT

dbSNP	Extend	SEQ ID	Term
rs#	Primer	NO.	Mix
775903	TCTGAGTGACAGAGCGA	<u>297</u>	ACT
776251	GCTCCCGCAACCTCCGC	<u>298</u>	ACG
892088	GCCTCGGCCGCAATCACA	<u>299</u>	ACT
892089	CGGTCACCGTGATGATGGG	300	ACT
892090	AGAATCCCGAGAGATGGTAC	<u>301</u>	CGT
892091	GTCCTTCACCTGAGCTTCC	<u>302</u>	ACT
1036231	GTGTTTACTGTTCAAGGCAAGT	<u>303</u>	ACT
1036232	GGCAACAGAGCAAGACT	<u>304</u>	ACG
1043673	ACTGAGAAACATCATCCCTGGG	<u>305</u>	CGT
1043678	AGTCACAGGCAGTTCACC	<u>306</u>	CGT
1043680	CTGTGACTCCTCTCCCC	<u>307</u>	ACT
1043684	CTGTTTTATACCTGCACAC	<u>308</u>	ACT
1054796	ACGCCAGGCAGGCTCTCA	<u>309</u>	ACT
1059211	CGCCTACTGCCAGAGCAAGCT	<u>310</u>	ACG
1064675	ATTCCTTTTTGCTGAAATAATGAA	<u>311</u>	ACT
1560714	TGGGGCGTGATGGCTCA	<u>312</u>	ACG
1613662	CAACAGAACCACCTTCC	<u>313</u>	ACG
1625609	GTGCACACCTGTAATCC	314	ACG
1625689	CAGGGCTCAAGCGATTCTCC	315	ACG
1626971	TCGCCTGGCCAAAAAA	316	ACT
1654406	CATTGTACCCCAGGTTGAAAAT	317	CGT
1654409	GTGGTACCACACCCAGCTAATT	318	ACT
1654410	ATCATACTGAGATGCTATCAGAA	319	ACT
1654411	GCACTTTGGGAGGTTGAGG	320	ACT
1654412	CATTCTCCCACCTCAGCCCCC	321	ACG
1654413	CCCGTTTGATTTCCGGGTC	322	CGT
1654415	GGCCCCATCACCCAAAA	323	ACG
1654416	GACCGTCTCATTCACAAAC	324	ACT
1654419	TTGGTGCTTCACTCTGAGAC	325	ACT
1654420	GAGAACCGCTGATCAATGCA	326	CGT
1654421	GCATGCAGCTCCCGTCC	327	ACT
1654424	CCACCTATGGCCGCGCCCCT	328	· ACT
1654425	CAGGGACCCATACCTGTGGTC	329	ACT
1654438	TCAGCCTCCTGAGTAGCTGG	330	ACT
1654439	GTTTCTGGAGCACTCCGGT	331	ACT
1654441	GATAAGCCTGGCCAACA	332	ACG
1654442	ATGATCTCGGCTCACTGCAA	333	ACT
1654444	TATGGATCTTTCTAGTCTTGTTT	334	CGT
1654446	ACTGATTACAGGCGTGC	335	ACT
1654447	CCCGATGCCTGTGTTGGC	336	ACT
1654448	AGTGCTGGGATTACAGG	337	ACG
1654449	AATCTCTCTTTTCCCTACACA	338	ACG
1654451	TAATGCGTACAGCAGCC	339	CGT
1654452	ACTGGAGGAGGATGCACTTA	340	ACG
	6	_ <del></del> _	Soria

dbSNP rs#	lbSNP Extend rs# Primer		Term Mix
	ATGCACAGAAACAAGGATCTA	NO.	ACT
1654459		341	ACT
1654485	CTTGCTTTTTTTTTTTTGGACAG	342	
1654491	GCACCCCGAGCCTTTCCAG	343	ACT
1654495	TTGTCGTAAGTCTCCTCTTT	344	CGT
1654496	CGGGAAGGTTGAAGTTGGAC	345	CGT
1654497	CCATTTACAACCAATTGC	346	ACT
1654498	CTTGTGGGACTTCTTTTTA	347	ACT
1654499	ACCCTGGCCTCCCTAAC	348	ACT
1654503	GGGCAACGTTTGAAGATGCTCTGC	<u>349</u>	ACG
1654504	CACCTTTGACTCTTGAGCC	<u>350</u>	ACT
1654505	TAGCTATGTGCCACCATGCC	<u>351</u>	ACG
1671133	GATTGTAGCTAACTCACAAGG	<u>352</u>	ACT
1671140	TACCTACCCACATCCTATAAAA	<u>353</u>	ACG
1671148	CCTGTAATCCCGGCACT	<u>354</u>	ACT
1671149	CTGGCCACAGTGGCTCA	<u>355</u>	ACG
1671150	CGGGTGACGAAGCCTGAC	<u>356</u>	ACG
1671151	TCCTCTGTGCAAAATCCTCC	<u>357</u>	ACG
1671152	CTCCATCCTGACCCCCGT	<u>358</u>	ACT
1671153	CTGTGGAATTGTGCCTC	<u>359</u>	CGT
1671169	CATGTCCCACAGAGGCTAAC	<u>360</u>	ACT
1671170	GAGAGGGCAATGCCTCAGAG	<u>361</u>	CGT
1671171	TTCTGGGATTCTCTAGAGGG	<u>362</u>	ACT
1671176	AGACATCATCACATCACACCA	<u>363</u>	CGT
1671178	CCAGGCTGGACTCGAACT	<u>364</u>	ACG
1671182	ACTGAATGTCTCGGTATAAAACC	<u>365</u>	ACG
1671187	CAGGTTGAGAAAGCTCTA	<u>366</u>	CGT
1671188	CAGAGTGAGGAGAGTGAGAC	367	ACT
1671191	GAGCGGTTAGAAGATGTGCT	368	ACT
1671192	AAGCCTGTAGGCTTTTAA	<u>369</u>	ACG
1671196	GGGATGACTGAATGAGACAGTA	370	ACG
1671198	CCCTCTTAGCAAATTTCAGCT	371	ACT
1671199	TAACTTTTTTGTGTGAGAA	372	ACG
1671214	CGTGCATCCTTCCCACCTA	373	ACG
1671215	GTACTCAAGATGATGTAA	374	CGT
1671216	TTACACCCTGGAGTGGTCC	375	ACT
1671217	TGCCCTCTCGCTGGCTGG	376	ACG
1671218	CAAAGGAGGTGGTCGCAC	377	ACG
1671219	CAGGAGAATGGTGTGAACC	378	ACG
1671221	AGCTGGGATTACAGGCA	379	ACG
1671223	TACAAAATTAGCTGGGCATG	380	ACT
1671224	CTGTGAGCCGAGATTGC	381	ACT
1671225	CTCAATGTGATCCTCCT	382	ACT
1671226	GCAGGAGAATCACTTGAACTT	383	ACT
1671227	AGATCATGCCATTGCCAGC	384	ACT

dbSNP rs#	Extend Primer	SEQ ID NO.	Term Mix
1671228	ACAGAAGTTAGCTGGGC	<u>385</u>	ACT
1869616	CTTCAACCTCCGCCTCCTGG	<u>386</u>	ACT
2019599	GAAAAGCATGGGCCGGGCA	<u>387</u>	ACG
2116883	CATACTCACCAATATCTGCT	<u>388</u>	ACT
2124090	GCTTTGTGTTCTTTCTAGTC	<u>389</u>	ACT
2163833	GCCAGCAATGCACGCGCAGT	<u>390</u>	ACG
2217659	GTAACTTACCTAATGATAGAGG	<u>391</u>	ACG
2304167	TGACTCCTTTGGACTGG	<u>392</u>	ACG
2304168	CAGTTCGGTGAAGTGGTT	<u>393</u>	ACT
2365593	GGTGTGAGCCACCACGCC	<u>394</u>	ACG
2365721	AGACTCCCTCTCAAAATAA	<u>395</u>	ACG
2569513	AGGGATAAGCATGAAACCACT	<u>396</u>	ACG
2569514	GCGTGAGCCACCACGCC	<u>397</u>	ACG
2886414	GGGTGACAAAGTGAGACTC	<u>398</u>	ACG
2886415	CACGCCTGGCTAAGCCT	<u>399</u>	ACT
3745912	GGCTGGCAGGCCAGGTCAAC	<u>400</u>	ACT
3786863	GTGCTGGGATTACAGGC	<u>401</u>	ACT

Please amend Table 11 on page 91 as follows:

**TABLE 11** 

dbSNP rs#	First PCR primer	SEQ ID NO.	Second PCR primer	SEQ ID
1654416	ACGTTGGATGTCTGCTGAGCATGAAATGCC		ACGTTGGATGCTGAACTGACCGTCTCATTC	

Please amend Table 12 on page 91 as follows:

TABLE 12

dbSNP	Extend  1. Primer	SEQ ID	Term
rs#		NO.	Mix
1654416	TGACCGTCTCATTCACAAAC	<u>404</u>	ACT

Please amend Table 15 on pages 94-95 as follows:

**TABLE 15** 

dbSNP rs#	Forward PCR primer	SEQ ID NO.	Reverse PCR primer	SEQ ID NO.
LAMA4_SNP5	ACGTTGGATGACAGTTCTTGGCTATCCTGG	<u>405</u>	ACGTTGGATGACTGGCCAGTGTAGGAATTG	<u>406</u>
LAMA4_SNP6	ACGTTGGATGGAAAGGGATTGACTCAGGAG	<u>407</u>	ACGTTGGATGCTTCCTTCACCTGAAGATGG	<u>408</u>

dbSNP rs#	Forward PCR primer	SEQ ID NO.	Reverse PCR primer	SEQ ID NO.
LAMA4_SNP4	ACGTTGGATGTTGAAGGACTGATCTATGGG	<u>409</u>	ACGTTGGATGAAAGCAACAGACAAGGCAAG	<u>410</u>
LAMA4_SNP1	ACGTTGGATGCAGACTGGAAATGCGCAATG	411	ACGTTGGATGCGTATCTTCAAGATGCACAG	<u>412</u>
1050348	ACGTTGGATGTGTTCATGTCTTCGGCATCC	<u>413</u>	ACGTTGGATGCAGCTGGATGACTACAATGC	<u>414</u>
LAMA4_SNP2	ACGTTGGATGAGGAATGCTTACAACGGAGG	<u>415</u>	ACGTTGGATGAACTCCCTTCATCCTTCCTC	<u>416</u>
744006	ACGTTGGATGTTGCCTTGAAGGTAGGCATG	<u>417</u>	ACGTTGGATGGGGTTAGCAGCTTAACTTTC	<u>418</u>
763247	ACGTTGGATGCCGGCCAAGACCAATACATC	<u>419</u>	ACGTTGGATGTGCAGACATGCACTATTCTC	420
764071	ACGTTGGATGCCACTTGGAAAGATTCAAGG	<u>421</u>	ACGTTGGATGTATTGTGACTTCTGCAGAAC	422
764587	ACGTTGGATGCAACATAGACCAGAAGTGGG	423	ACGTTGGATGTTCACATACGGAAGGCCTTG	424
969138	ACGTTGGATGACTGGACCAAGGTAGATCAC	<u>425</u>	ACGTTGGATGCTCAGGCTAATCTCTCTAGG	<u>426</u>
971402	ACGTTGGATGCCACTTTTCTGTGGAAATATC	<u>427</u>	ACGTTGGATGCAAGTTAATGAGTTTCTCCC	<u>428</u>
971405	ACGTTGGATGAAACAGTGCTTTTGAAGGAG	<u>429</u>	ACGTTGGATGCTATCTCCAAAGGGTAACAG	<u>430</u>
1050348	ACGTTGGATGCAGCTGGATGACTACAATGC	<u>431</u>	ACGTTGGATGTGTTCATGTCTTCGGCATCC	<u>432</u>
1050349	ACGTTGGATGCTATGATTTTGGATTCAGCG	<u>433</u>	ACGTTGGATGACCTCATGGTATTTTGCATC	434
1158747	ACGTTGGATGTTGAAGGACTGATCTATGGG	435	ACGTTGGATGAAAGCAACAGACAAGGCAAG	436
1418499	ACGTTGGATGACCATAGGGAACTAGAAATC	437	ACGTTGGATGCTTTAAGATAGATTCCCAGGG	<u>438</u>
1480646	ACGTTGGATGCAGTGTCTCTTCCTTTCCAG	439	ACGTTGGATGCAAATTTCCACGAGCCTGAG	<u>440</u>
1894681	ACGTTGGATGTGGGATTCCCCTAAAGGATG	441	ACGTTGGATGAAGATCAGCAGCACCAAAGG	442
2032565	ACGTTGGATGAAAGAGCAACTGAAGGACCC	443	ACGTTGGATGTAAATTGGAACATCAACAGG	444
2032566	ACGTTGGATGTAAATTGGAACATCAACAGG	445	ACGTTGGATGAAAGAGCAACTGAAGGACCC	446
2032567	ACGTTGGATGCGTATCTTCAAGATGCACAG	447	ACGTTGGATGAGACTGGAAATGCGCAATGG	448
2032568	ACGTTGGATGACTCGCATAACAGATGTTCC	449	ACGTTGGATGTAACCATTGCGCATTTCCAG	<u>450</u>
2051649	ACGTTGGATGACCTGCTGAAAACCAACACC	451	ACGTTGGATGGGAGAGGAGAACCCTGGAC	<u>452</u>
2068770	ACGTTGGATGCACTTCACGTACTTCACTGG	<u>453</u>	ACGTTGGATGAGTTTGCTCCTATGTGGCTC	<u>454</u>
2072019	ACGTTGGATGAGGTCCACAGAAGATGTTAG	<u>455</u>	ACGTTGGATGCACAACGGTCATTTGAACAC	<u>456</u>
2072020	ACGTTGGATGAAGTCCTGTTGTCTGCAAGG	<u>457</u>	ACGTTGGATGCAGTTGTCTTAGCACACAGG	<u>458</u>
2072022	ACGTTGGATGCAAAGAAGAAGATGTAGTGG	<u>459</u>	ACGTTGGATGCGAAATCTGGTCCTATGAAG	<u>460</u>
2072026	ACGTTGGATGTCCTATCACCATCACACTAC	<u>461</u>	ACGTTGGATGCAGCATCAAACAGAATAGGC	<u>462</u>
2072029	ACGTTGGATGTCCTTGCAGACTGATACTCC	<u>463</u>	ACGTTGGATGTCACTCACTCCTTGCTAAGC	<u>464</u>
2157544	ACGTTGGATGCATATGTAGTAGGAATGAGGG	<u>465</u>	ACGTTGGATGTGAGGCTCAAAGGGATTAGG	<u>466</u>
2157545	ACGTTGGATGTCTGGTCAACCACATAGATC	<u>467</u>	ACGTTGGATGTGTTCTACTGCAGCTCCAAG	<u>468</u>
2157546	ACGTTGGATGTCCACTTGTACAGAATGGAG	<u>469</u>	ACGTTGGATGCATTTACTCAGTGCCAGGTC	<u>470</u>
2157547	ACGTTGGATGCCATACCATTTACTTCTGCC	<u>471</u>	ACGTTGGATGAGGCAAGTACACATACAATG	<u>472</u>
2157550	ACGTTGGATGCACACACATTTTAATTGCC	<u>473</u>	ACGTTGGATGTTGTTCAGAATTACATGATG	<u>474</u>
2213838	ACGTTGGATGAAAGGACTTGAGGGTGATTG	<u>475</u>	ACGTTGGATGGCAACAAACAGTGTTCCAGC	<u>476</u>
2213839	ACGTTGGATGAGTCACAGTTCAGTCCCAAC	<u>477</u>	ACGTTGGATGGGGCAATTTTCTAGTCCAGC	<u>478</u>
2213840	ACGTTGGATGCTTTCGCACAAGGCTCTATC	<u>479</u>	ACGTTGGATGAAGTCTGTGTTTAAGCCCCC	<u>480</u>
2227237	ACGTTGGATGGATGTCTCTAAGTTGAAATGC	481	ACGTTGGATGATATCAATCACCCTCAAGTC	<u>482</u>
2237238	ACGTTGGATGCAGAGGCTGAAGGAACATAC	<u>483</u>	ACGTTGGATGTCTGTAATCCCAGGACCCTA	<u>484</u>
2237241	ACGTTGGATGTCAGCAGGGCTCTATCTAAG	<u>485</u>	ACGTTGGATGCCAAGCAGTATTGCTAATGG	<u>486</u>
2237242	ACGTTGGATGCCTCACCATTGTGTTTAGGC	<u>487</u>	ACGTTGGATGTGACTATTTCCGCTTGGCTC	488
2237244	ACGTTGGATGGAGAAAAATAGACTCGGCCC	<u>489</u>	ACGTTGGATGCACAGACGCAGGATTTGGAT	490
2237247	ACGTTGGATGCTGCTTCTCCAGTAATGTTG	<u>491</u>	ACGTTGGATGGTGTTAGTAACACTGATGCC	492
2237248	ACGTTGGATGCCCTCCCCAGATATCATTAG	<u>493</u>	ACGTTGGATGCATATCCACAGCCTAATCAC	494
2237249	ACGTTGGATGAAATGCTTCCTACTGCAATC	<u>495</u>	ACGTTGGATGTGGAGAGTTGTGGTTGATGG	<u>496</u>
2239849	ACGTTGGATGGACATCAGATCAGACAGCAC	<u>497</u>	ACGTTGGATGACTTTCTGGCATTGACTGGG	<u>498</u>
2239850	ACGTTGGATGGCCCAGGAAAAATTAATTCAC	<u>499</u>	ACGTTGGATGGCAGTACGGATTAGCATGAG	<u>500</u>

dbSNP rs#	Forward PCR primer	SEQ ID NO.	Reverse PCR primer	SEQ ID NO.
2269646	ACGTTGGATGTCACCTCACTTTTGAAGAGC	501	ACGTTGGATGTCTGGTTAGGCTTCAGTTAG	502
2277084	ACGTTGGATGGAGGGTAAAAATGACAGCAG	503	ACGTTGGATGTTTTGCTTGGTGTTTAGCAG	504
2282853	ACGTTGGATGTCTTGACCTTCCTGGTTTTC	<u>505</u>	ACGTTGGATGTATCAGAGCTAGAAGAAACC	506
2282854	ACGTTGGATGTAGCCAGTGGTTAAGAAAGC	507	ACGTTGGATGTTCTCATGTTGGGGAGACAC	<u>508</u>
2301512	ACGTTGGATGATCTGAGTGGTTTCAGGAGG	<u>509</u>	ACGTTGGATGACCTGTTGGAACACATGAAG	<u>510</u>
2301513	ACGTTGGATGCTGGCGGGTAGTGTCTTCAT	<u>511</u>	ACGTTGGATGCTTTGAAATTGTTCTTGTCC	<u>512</u>
2345808	ACGTTGGATGTTCTGGGATTTAAAGGAGGC	<u>513</u>	ACGTTGGATGCCAAACATTTCTTGTTGGAC	<u>514</u>
3734286	ACGTTGGATGACCTTACACTCCAGTGAATC	<u>515</u>	ACGTTGGATGGCCGTTAAGCAACTACAAGC	<u>516</u>
3734287	ACGTTGGATGCAGTGGAGAAGATGAAACCC	<u>517</u>	ACGTTGGATGCCCACTTCTGGTCTATGTTG	<u>518</u>
3752577	ACGTTGGATGCATGGCTGAGGTTACTTAGG	<u>519</u>	ACGTTGGATGGAATGCGTCAGGGATTTATG	<u>520</u>
3777925	ACGTTGGATGCTACAAGTCTAACAGTCAGAG	<u>521</u>	ACGTTGGATGTTACAGAGCAAGGTCTGAGG	<u>522</u>
3777926	ACGTTGGATGGTGAGTACCATCCCTTTTGC	<u>523</u>	ACGTTGGATGCTGTTAAACTGCCTCAGACC	<u>524</u>
3777927	ACGTTGGATGAAACGAATGCTTGAGAGCAG	<u>525</u>	ACGTTGGATGGTCCTGATTTATGAGCTCCC	<u>526</u>
3777928	ACGTTGGATGTTCACACGTAGACCCTGTTG	<u>527</u>	ACGTTGGATGTCAGGAGTTGAGCAAGCTAG	<u>528</u>
3777929	ACGTTGGATGGCTGTCTTTGGGATTAAAT	<u>529</u>	ACGTTGGATGTTCATAAAGAAGTGGAGAGC	<u>530</u>
3777932	ACGTTGGATGTCCCAGACCTTAAGATTCCC	<u>531</u>	ACGTTGGATGTATTAGGCTCTTTGGCCGAC	<u>532</u>
3777934	ACGTTGGATGCAAGATCCAGATGGTGAGGG	<u>533</u>	ACGTTGGATGCAAGGTCAGAGTGTCACTGG	<u>534</u>
3777941	ACGTTGGATGGCTTCCTGAGATTATATTGAC	<u>535</u>	ACGTTGGATGCTCCATTCCAAATTCCTTTC	<u>536</u>
3777942	ACGTTGGATGTCATGACAAATCATGACTAG	<u>537</u>	ACGTTGGATGTCAGATACAAGTGAAGGTAG	<u>538</u>
3798357	ACGTTGGATGTCCCAATTCAGGAAATGGTG	<u>539</u>	ACGTTGGATGTGCTTGGTATACCATGCCTG	<u>540</u>
3798359	ACGTTGGATGTTCCTCAGCACACAGCCCCA	<u>541</u>	ACGTTGGATGATGAACCTTACACAGGCCAG	<u>542</u>
3822941	ACGTTGGATGTATAATAAACTGATAGTTGC	<u>543</u>	ACGTTGGATGCTCTGTACTTAGGACACACG	<u>544</u>
3948760	ACGTTGGATGTCCACTTGTACAGAATGGAG	<u>545</u>	ACGTTGGATGTCCCACACTCAAAACTTTGC	<u>546</u>

Please amend Table 16 on pages 95-97 as follows:

TABLE 16

dbSNP rs#	Extend Primer	SEQ ID NO.	Term Mix
LAMA4_SNP5	ATTGCTTACGCAACACCAC	<u>547</u>	ACG
LAMA4_SNP6	AGATGGAGAGAATGCCAC	<u>548</u>	CGT
LAMA4_SNP4	GCAAGTGGGCATTCGACCA	<u>549</u>	CGT
LAMA4_SNP1	CTTCAAGATGCACAGGGCCAC	<u>550</u>	ACG
1050348	CACTTGACCAGGCCCTTAAC	<u>551</u>	ACG
LAMA4_SNP2	GGCCCGCCTGCATCTGTG	<u>552</u>	ACG
744006	CTTTCTCTCTTTCCAGG	<u>553</u>	ACG
763247	CTTTAATCCCCCACACT	<u>554</u>	ACT
764071	AGAACATATATGTTGCATTTTTT	<u>555</u>	ACT
764587	GAAGGCCTTGCCTGTTA	<u>556</u>	ACT
969138	AGGAAGAGAATCTGATAGCC	<u>557</u>	ACT
971402	AGTITCTCCCACTTACC	<u>558</u>	ACT
971405	AGGGTAACAGAATGATTAAAA	<u>559</u>	CGT

1050348         TTCGGCATCCCTGACAT         560         ACT           1050349         CGTATCTTCAAGATGCACA         561         ACT           1158747         GCAAGTGGGCATTCGACCA         562         CGT           1418499         GGGCAGAATTACTGAATCAAG         563         ACT           1480646         CAGCAGACTCTGATGTGGC         564         ACC           1894681         GGGAGCATCTTTTGAGC         565         ACT           2032565         CAACAGGAAAAATACATCCA         566         CGT           2032566         CAACCCTAGGAAAACATTCAGC         568         ACT           2032567         TTCTATGATTTTGGATTCAGC         568         ACT           2032568         ACATACTCTGAGGAGAGAAAG         569         ACT           2051649         GAACCCTGGACAAGAAT         570         ACT           2078019         TTTGAACACTACCAGTTTCTGTAT         572         ACT           2072020         AAACAATCCATTTAACATACCTA         573         ACC           2072022         GCAAATGATTCTGAGGTGTT         575         ACC           2072029         CCTGGCAATGATCAACCCC         576         ACT           2157544         CTAAATATTAGCAGACTGAAATAC         577         ACT           2157545         GCTGGCATAAA	dbSNP	Extend	SEQ ID	Term
1050349         CGTATCTTCAAGATGCACA         561         ACT           1158747         GCAAGTGGGCATTCGACCA         562         CGT           1418499         GGGCAGAATTACTGAATCAAG         563         ACT           1480646         CAGCAGACTCTGATGTGGC         564         ACC           1894681         GGGAGCATCTTTTGAGC         565         ACT           2032565         CAACAGGAAAAATACATCCA         566         CGT           2032566         CAACCCTAGGAAAACATTT         567         ACT           2032567         TTCTATGATTTTGGATTCAGC         568         ACT           2032568         ACATACTCTGAGGAGAAGAAAG         569         ACT           2032569         GAACCCTGGACAAGAAT         570         ACT           2051649         GAACCCTGGACAAGAAT         570         ACT           2072019         TTGAACACTACAGTTTCTGTAT         572         ACT           2072020         AAACAATCCATTTAACATACCTA         573         ACC           2072022         GCAAATGAATTCTGGGA         574         ACT           2072029         CCTGGCAATGATCAACCCCC         576         ACT           2157544         CTAAATATTAGCAGACTGAAATAC         572         ACT           2157545         GTGCCAGGTCCCACACT				Mix
1158747         GCAAGTGGGCATTCGACCA         562         CGT           1418499         GGGCAGAATTACTGAATCAAG         563         ACT           1480646         CAGCAGACTCTGATGTGGC         564         ACC           1894681         GGGAGCATCTTTTGAGC         565         ACT           2032565         CAACAGGAAAAATACATCCA         566         CGT           2032566         CAACCCTAGGAAAACATTT         567         ACT           2032568         ACATACTCTGAGGAGAGAAAG         569         ACT           2051649         GAACCCTGGACAAGAAT         570         ACT           2072019         TTTGAACACTACAGTTTCTGTTAT         572         ACT           2072020         AAACAATCCATTTAACATACCTA         573         ACC           2072022         GCAAATGAATTCTGGGA         574         ACT           2072022         GCAAATGAATCTTTGAGGTGTT         575         ACT           2072029         CCTGGCAATGATCAACCCCC         576         ACT           2157544         CTAAATATTAGCAGACTGAAATAC         577         ACT           2157545         GCTGGCATAAATGAAATTTATAGAATT         578         ACT           2157546         GTGCCAGGTCCCACACT         579         ACT           2157550         TTACATG	1050348	TTCGGCATCCCTGACAT		ACT
1418499         GGGCAGAATTACTGAATCAAG         563         ACT           1480646         CAGCAGACTCTGATGTGGC         564         ACC           1894681         GGGAGCATCTTTTGAGC         565         ACT           2032565         CAACAGGAAAAATACATCCA         566         CGT           2032566         CAACCCTAGGAAAACATTT         567         ACT           2032568         ACATACTCTGAGGAGAGAAAG         569         ACT           2051649         GAACCCTGGACAAGAAT         570         ACT           2051649         GAACCCTGGACAAGAAT         570         ACT           2072019         TTTGAACACTACAGTTTCTGTTAT         572         ACT           2072020         AAACAATCCATTTAACATACCTA         573         ACC           2072022         GCAAATGAATTCTGGGA         574         ACT           2072029         CCTGGCAATGATCAACCCCC         576         ACT           2157544         CTAAATATTAGCAGACTGAAATAC         577         ACT           2157545         GCTGGCAATGAATGAATTTATC         580         ACT           2157546         GTGCCAGGTCCCACACT         579         ACT           2157550         TTACATGATGAATTTATGGAAGT         581         ACT           2213838         TTCCAGCATGATA	1050349	CGTATCTTCAAGATGCACA	<u>561</u>	ACT
1480646         CAGCAGACTCTGATGTGGC         564         ACC           1894681         GGGAGCATCTTTTGAGC         565         ACT           2032565         CAACAGGAAAAATACATCCA         566         CG           2032566         CAACCCTAGGAAAACATTT         567         ACT           2032567         TTCTATGATTTTGGATTCAGC         568         ACT           2032568         ACATACTCTGAGGAGAGAAAG         569         ACT           2051649         GAACCCTGGACAAGAAT         570         ACT           2068770         ATGTGGCTCAAACATCCGAA         571         ACT           2072019         TTTGAACACTACAGTTTCTGTTAT         572         ACT           2072020         AAACAATCCATTTAACATACCTA         573         ACC           2072022         GCAAATGAATTCTGGGA         574         ACT           2072029         CCTGGCAATGATCAACCCCC         576         ACT           2157544         CTAAATATTAGCAGACTGAAATAC         577         ACT           2157545         GCTGGCATAAATGAATTTATGGAAGT         578         ACC           2157546         GTGCCAGGTCCCACACT         579         ACT           2157547         GTACACATGAATATTATGGAAGT         581         ACT           2157550         TTACATG	1158747	GCAAGTGGGCATTCGACCA	<u>562</u>	CGT
1894681         GGGAGCATCTTTTGAGC         565         ACT           2032565         CAACAGGAAAAATACATCCA         566         CG           2032566         CAACCCTAGGAAAACATTT         567         ACT           2032567         TTCTATGATTTTGGATTCAGC         568         ACT           2032568         ACATACTCTGAGGAGAGAAAG         569         ACT           2051649         GAACCCTGGACAAGAAT         570         ACT           2072019         TTTGAACACTACAGTTTCTGTTAT         572         ACT           2072020         AAACAATCCATTTAACATACCTA         573         ACC           2072021         GCAAATGAATTCTGGGA         574         ACT           2072022         GCAAATGAATTCTGGGA         574         ACT           2072029         CCTGGCAATGATCAACCCCC         576         ACT           2157544         CTAAATATTAGCAGACTGAAATAC         577         ACT           2157545         GCTGGCATAAATGAATTTACTC         580         ACT           2157546         GTGCCAGGTCCCACACT         579         ACT           2157547         GTACACATACAATGAATTTACTC         580         ACT           2157550         TTACATGATGAATATTATGGAACA         581         ACT           2213838         TTCCAGCAT	1418499	GGGCAGAATTACTGAATCAAG	<u>563</u>	ACT
2032565         CAACAGGAAAAATACATCCA         566         CG           2032566         CAACCCTAGGAAAACATTT         567         ACT           2032567         TTCTATGATITTGGATTCAGC         568         ACT           2032568         ACATACTCTGAGGAGAAAAG         569         ACT           2051649         GAACCCTGGACAAGAAT         570         ACT           2068770         ATGTGGCTCAAACATCCGAA         571         ACT           2072019         TTTGAACACTACAGTITCTGTTAT         572         ACT           2072020         AAACAATCCATTTAACATACCTA         573         ACC           2072022         GCAAATGAATTCTGGGA         574         ACT           2072026         TGAAAGTCTTTGAGGTGTT         575         ACC           2072029         CCTGGCAATGATCAACCCCC         576         ACT           2157544         CTAAATATTAGCAGACTGAAATAC         577         ACT           2157545         GCTGGCATAAATGAAATTATTAC         578         ACC           2157546         GTGCCAGGTCCCACACT         579         ACT           2157547         GTACACATACAATGATTTACTC         580         ACT           2157550         TTACATGAGATACAGTAAAATT         581         ACT           2213838         TCCAGGA	1480646	CAGCAGACTCTGATGTGGC	<u>564</u>	ACG
2032566         CAACCCTAGGAAAACATTT         567         ACT           2032567         TTCTATGATTTTGGATTCAGC         568         ACT           2032568         ACATACTCTGAGGAGAGAAAG         569         ACT           2051649         GAACCCTGGACAAGAAT         570         ACT           2068770         ATGTGGCTCAAACATCCGAA         571         ACT           2072019         TTTGAACACTACAGTTTCTGTTAT         572         ACT           2072020         AAACAATCCATTTAACATACCTA         573         ACC           2072022         GCAAATGAATTCTGGGA         574         ACT           2072029         CCTGGCAATGATCAACCCCC         576         ACT           2072029         CCTGGCAATGATCAACCCCC         576         ACT           2157544         CTAAATATTAGCAGACTGAAATAC         577         ACT           2157545         GCTGGCATAAATGAAATTA         578         ACC           2157546         GTGCCAGGTCCCACACT         579         ACT           2157547         GTACACATGAATATTATGGAAGT         581         ACT           2157550         TTACATGATGATATATTATGGAAGT         581         ACT           2213838         TTCCAGCATGATTCTAAGACA         582         ACT           2213840         TGA	1894681	GGGAGCATCTTTTGAGC	<u>565</u>	ACT
2032567         TTCTATGATTITIGGATTCAGC         568         ACT           2032568         ACATACTCTGAGGAGAGAAAG         569         ACT           2051649         GAACCCTGGACAAGAAT         570         ACT           2068770         ATGTGGCTCAAACATCCGAA         571         ACT           2072019         TTTGAACACTACAGTTTCTGTTAT         572         ACT           2072020         AAACAATCCATTTAACATACCTA         573         ACC           2072022         GCAAATGAATTCTGGGA         574         ACT           2072026         TGAAAGTCTTTGAGGTGTT         575         ACC           2072029         CCTGGCAATGATCAACCCCC         576         ACT           2157544         CTAAATATTAGCAGACTGAAATAC         577         ACT           2157545         GCTGGCATAAATGAAATTG         578         ACC           2157546         GTGCCAGGTCCCACACT         579         ACT           2157547         GTACACATACAATGATTTTACTC         580         ACT           2157550         TTACATGATGAATATTATGGAAGT         581         ACT           2213838         TTCCAGCATGATTCTAAGACA         582         ACT           22313840         TGAAATGAATCTCCCAATAGAC         584         ACC           22372241 <td< td=""><td>2032565</td><td>CAACAGGAAAAATACATCCA</td><td><u>566</u></td><td>CGT</td></td<>	2032565	CAACAGGAAAAATACATCCA	<u>566</u>	CGT
2032568         ACATACTCTGAGGAGAGAAAG         569         ACT           2051649         GAACCCTGGACAAGAAT         570         ACT           2068770         ATGTGGCTCAAACATCCGAA         571         ACT           2072019         TTTGAACACTACAGTTTCTGTTAT         572         ACT           2072020         AAACAATCCATTTAACATACCTA         573         ACC           2072022         GCAAATGAATTCTGGGA         574         ACT           2072026         TGAAAGTCTTTGAGGTGTT         575         ACC           2072029         CCTGGCAATGATCAACCCCC         576         ACT           2157544         CTAAATATTAGCAGACTGAAATAC         577         ACT           2157545         GCTGGCATAAATGAAATTG         578         ACC           2157546         GTGCCAGGTCCCACACT         579         ACT           2157547         GTACACATACAATGATTTTACTC         580         ACT           2157550         TTACATGATGAATATTATGGAAGT         581         ACT           2213838         TTCCAGCATGATTCTAAGACA         582         ACT           2213840         TGAAATGAATTCTCCAATAGAC         584         ACC           2237238         TCCCAGGACCCTAAAAAAGT         586         CG           2237241         CAGT	2032566	CAACCCTAGGAAAACATTT	<u>567</u>	ACT
2051649         GAACCCTGGACAAGAAT         570         ACT           2068770         ATGTGGCTCAAACATCCGAA         571         ACT           2072019         TITGAACACTACAGTITCTGTTAT         572         ACT           2072020         AAACAATCCATTTAACATACCTA         573         ACC           2072022         GCAAATGAATTCTGGGA         574         ACT           2072029         CCTGGCAATGATCAACCCCC         576         ACT           2157544         CTAAATATTAGCAGACTGAAATAC         577         ACT           2157545         GCTGGCATAAATGAAATTG         578         ACC           2157546         GTGCCAGGTCCCACACT         579         ACT           2157547         GTACACATACAATGATTTTACTC         580         ACT           2157550         TTACATGATGAATATTATGGAAGT         581         ACT           2213838         TTCCAGCATGATTCTAAGACA         582         ACT           2213840         TGAAATGAATTCTCCAATAGAC         584         ACC           2227237         ACCCTCAAGTCCTTTTG         585         ACC           2237241         CAGTATTGCTAATGGGTGTTC         587         ACT           2237242         TGTCTCTAGGGCACTACATATC         588         ACT           2237244         GAA	2032567	TTCTATGATTTTGGATTCAGC	<u>568</u>	ACT_
2068770         ATGTGGCTCAAACATCCGAA         571         ACT           2072019         TITGAACACTACAGTITCTGTTAT         572         ACT           2072020         AAACAATCCATTTAACATACCTA         573         ACC           2072022         GCAAATGAATTCTGGGA         574         ACT           2072029         CCTGGCAATGATCAACCCCC         576         ACT           2157544         CTAAATATTAGCAGACTGAAATAC         577         ACT           2157545         GCTGGCATAAATGAAATTG         578         ACC           2157546         GTGCCAGGTCCCACACT         579         ACT           2157550         TTACATGATGAATATTATGGAAGT         581         ACT           2157550         TTACATGATGAATATTATGGAAGT         581         ACT           2213838         TTCCAGCATGATTCTAAGACA         582         ACT           2213840         TGAAATGAATACAGTAAAAAATT         583         ACC           2237237         ACCCTCAAGTCCTTTTG         585         ACC           2237241         CAGTATTGCTAATAGACT         587         ACT           2237242         TGTCTCTAGGGCACTACATATC         588         ACT           2237244         GAAATAATGCTTCAGGGG         589         ACT           2237247         ATG	2032568	ACATACTCTGAGGAGAGAAAG	<u>569</u>	ACT
2072019         TTTGAACACTACAGTITCTGTTAT         572         ACT           2072020         AAACAATCCATTTAACATACCTA         573         ACC           2072022         GCAAATGAATTCTGGGA         574         ACT           2072026         TGAAAGTCTTTGAGGTGTT         575         ACC           2072029         CCTGGCAATGATCAACCCCC         576         ACT           2157544         CTAAATATTAGCAGACTGAAATAC         577         ACT           2157545         GCTGGCATAAATGAATTG         578         ACC           2157546         GTGCCAGGTCCCACACT         579         ACT           2157547         GTACACATACAATGATTTTACTC         580         ACT           2157550         TTACATGATGAATATTATGGAAGT         581         ACT           2213838         TTCCAGCATGATTCTAAGACA         582         ACT           2213840         TGAAATGAATTCTCCAATAGAC         584         ACC           2227237         ACCCTCAAGTCCTTTTG         585         ACC           2237241         CAGTATTGCTAATGGGTGTTC         587         ACT           2237242         TGTCTCTAGGGCACTACATATC         588         ACT           2237244         GAAATAATGCTTCAGGGG         589         ACT           2237247         ATGC	2051649	GAACCCTGGACAAGAAT	<u>570</u>	ACT
2072020         AAACAATCCATTTAACATACCTA         573         ACC           2072022         GCAAATGAATTCTGGGA         574         ACT           2072026         TGAAAGTCTTTGAGGTGTT         575         ACC           2072029         CCTGGCAATGATCAACCCCC         576         ACT           2157544         CTAAATATTAGCAGACTGAAATAC         577         ACT           2157545         GCTGGCATAAATGAAATTG         578         ACC           2157546         GTGCCAGGTCCCACACT         579         ACT           2157550         GTACACATACAATGATTTATGGAAGT         581         ACT           2213838         TTCCAGCATGATTCTAAGACA         582         ACT           2213840         TGAAATGAATTCTCCAATAGAC         584         ACC           2227237         ACCCTCAAGTCCTTTTG         585         ACC           2237238         TCCCAGGACCCTAAAAAAAGT         586         CG           2237241         CAGTATTGCTAATGGGTGTTC         587         ACT           2237242         TGTCTCTAGGGCACTACATATC         588         ACT           2237244         GAAATAATGCTTCAGGGG         589         ACT           2237247         ATGCCTTCTAATGCATTCATTTTA         590         ACC           2237248         CCTA	2068770	ATGTGGCTCAAACATCCGAA	<u>571</u>	·ACT
2072022         GCAAATGAATTCTGGGA         574         ACT           2072026         TGAAAGTCTTTGAGGTGTT         575         ACC           2072029         CCTGGCAATGATCAACCCCC         576         ACT           2157544         CTAAATATTAGCAGACTGAAATAC         577         ACT           2157545         GCTGGCATAAATGAAATTG         578         ACT           2157546         GTGCCAGGTCCCACACT         579         ACT           2157557         GTACACATACAATGATTTACTC         580         ACT           2157550         TTACATGATGAATATTATGGAAGT         581         ACT           2213838         TTCCAGCATGATTCTAAGACA         582         ACT           2213840         TGAAATGAATTCTCCAATAGAC         584         ACC           2227237         ACCCTCAAGTCCTTTTG         585         ACC           2237238         TCCCAGGACCCTAAAAAAAGT         586         CG           2237241         CAGTATTGCTAATGGGTGTTC         587         ACT           2237242         TGTCTCTAGGGCACTACATATC         588         ACT           2237244         GAAATAATGCTTCAGGGG         589         ACT           2237247         ATGCCTTCTAATGCATTCATTITA         590         ACC           2237248         CCTAATC	2072019	TTTGAACACTACAGTTTCTGTTAT	<u>572</u>	ACT
2072026         TGAAAGTCTTTGAGGTGTT         575         ACC           2072029         CCTGGCAATGATCAACCCCC         576         ACT           2157544         CTAAATATTAGCAGACTGAAATAC         577         ACT           2157545         GCTGGCATAAATGAAATTG         578         ACC           2157546         GTGCCAGGTCCCACACT         579         ACT           2157550         GTACACATACAATGATTTTACTC         580         ACT           2213838         TTCCAGCATGATTCTAAGACA         582         ACT           2213839         CAACTTGAGATACAGTAAAAATT         583         ACC           2213840         TGAAATGAATTCTCCAATAGAC         584         ACC           2227237         ACCCTCAAGTCCTTTTG         585         ACC           2237238         TCCCAGGACCCTAAAAAAGT         586         CG           2237241         CAGTATTGCTAATGGGTGTTC         587         ACT           2237242         TGTCTCTAGGGCACTACATATC         588         ACT           2237244         GAAATAATGCTTCATGGGG         589         ACT           2237248         CCTAATCACATAAACCAGGAA         591         ACC           2237249         GAAAACAAGAGAGAGGGAAG         592         ACT           2239849         TGTGACT	2072020	AAACAATCCATTTAACATACCTA	<u>573</u>	ACG
2072029         CCTGGCAATGATCAACCCCC         576         ACT           2157544         CTAAATATTAGCAGACTGAAATAC         577         ACT           2157545         GCTGGCATAAATGAAATTG         578         ACC           2157546         GTGCCAGGTCCCACACT         579         ACT           2157547         GTACACATACAATGATTTTACTC         580         ACT           2157550         TTACATGATGAATATTATGGAAGT         581         ACT           2213838         TTCCAGCATGATTCTAAGACA         582         ACT           2213840         TGAAATGAGTACAGTAAAAATT         583         ACC           2227237         ACCCTCAAGTCCTTTTG         585         ACC           2237238         TCCCAGGACCCTAAAAAAGT         586         CG           2237241         CAGTATTGCTAATGGGTGTTC         587         ACT           2237242         TGTCTCTAGGGCACTACATATC         588         ACT           2237244         GAAATAATGCTTCAGGGG         589         ACT           2237247         ATGCCTTCTAATGCATTCATTTTA         590         ACC           2237248         CCTAATCACATAAACCAGGAA         591         ACC           2237249         GAAAACAAGAGAGAGGGAAG         592         ACT           2239849         TG	2072022	GCAAATGAATTCTGGGA	<u>574</u>	ACT
2157544         CTAAATATTAGCAGACTGAAATAC         577         ACT           2157545         GCTGGCATAAATGAAATTG         578         ACC           2157546         GTGCCAGGTCCCACACT         579         ACT           2157547         GTACACATACAATGATTTTACTC         580         ACT           2157550         TTACATGATGAATATTATGGAAGT         581         ACT           2213838         TTCCAGCATGATTCTAAGACA         582         ACT           2213840         TGAAATGAATTCTCCAATAGAC         584         ACC           2227237         ACCCTCAAGTCCTTTTG         585         ACC           2237238         TCCCAGGACCCTAAAAAAGT         586         CG           2237241         CAGTATTGCTAATGGGTGTTC         587         ACT           2237242         TGTCTCTAGGGCACTACATATC         588         ACT           2237244         GAAATAATGCTTCAGGGG         589         ACT           2237247         ATGCCTTCTAATGCATTCATTITA         590         ACC           2237248         CCTAATCACATAAACCAGGAA         591         ACC           2237249         GAAAACAAGAGAGAGGGAAG         592         ACT           2239849         TGTGACTCCTCATGCTAATC         593         ACC           2239850         CC	2072026	TGAAAGTCTTTGAGGTGTT	<u>575</u>	ACG
2157545         GCTGGCATAAATGAAATTG         578         ACC           2157546         GTGCCAGGTCCCACACT         579         ACT           2157547         GTACACATACAATGATTTACTC         580         ACT           2157550         TTACATGATGAATATTATGGAAGT         581         ACT           2213838         TTCCAGCATGATTCTAAGACA         582         ACT           2213849         CAACTTGAGATACAGTAAAAATT         583         ACC           2227237         ACCCTCAAGTCCTTTTG         585         ACC           2237238         TCCCAGGACCCTAAAAAAAGT         586         CGT           2237241         CAGTATTGCTAATGGGTGTTC         587         ACT           2237242         TGTCTCTAGGGCACTACATATC         588         ACT           2237244         GAAATAATGCTTCAGGGG         589         ACT           2237247         ATGCCTTCTAATGCATTCATTTTA         590         ACC           2237248         CCTAATCACATAAACCAGGAA         591         ACC           2237249         GAAAACAAGAGAGGGGAAG         592         ACT           2239849         TGTGACTCCTCATGCTAATC         593         ACC           2239850         CCAGTCAATGCCAGAAA         594         ACT	2072029	CCTGGCAATGATCAACCCCC	<u>576</u>	ACT
2157546         GTGCCAGGTCCCACACT         579         ACT           2157547         GTACACATACAATGATTTTACTC         580         ACT           2157550         TTACATGATGAATATTATGGAAGT         581         ACT           2213838         TTCCAGCATGATTCTAAGACA         582         ACT           2213839         CAACTTGAGATACAGTAAAAATT         583         ACC           2213840         TGAAATGAATTCTCCAATAGAC         584         ACC           2227237         ACCCTCAAGTCCTTTTG         585         ACC           2237238         TCCCAGGACCCTAAAAAAGT         586         CG           2237241         CAGTATTGCTAATGGGTGTTC         587         ACT           2237242         TGTCTCTAGGGCACTACATATC         588         ACT           2237244         GAAATAATGCTTCAGGGG         589         ACT           2237247         ATGCCTTCTAATGCATTCATTTTA         590         ACC           2237248         CCTAATCACATAAACCAGGAA         591         ACC           2237249         GAAAACAAGAGAGAGGGAAG         592         ACT           2239849         TGTGACTCCTCATGCTAATC         593         ACC           2239850         CCAGTCAATGCCAGAAA         594         ACT	2157544	CTAAATATTAGCAGACTGAAATAC	<u>577</u>	ACT
2157547         GTACACATACAATGATTTTACTC         580         ACT           2157550         TTACATGATGAATATTATGGAAGT         581         ACT           2213838         TTCCAGCATGATTCTAAGACA         582         ACT           2213839         CAACTTGAGATACAGTAAAAATT         583         ACC           2213840         TGAAATGAATTCTCCAATAGAC         584         ACC           2227237         ACCCTCAAGTCCTTTTG         585         ACC           2237238         TCCCAGGACCCTAAAAAAGT         586         CG           2237241         CAGTATTGCTAATGGGTGTTC         587         ACT           2237242         TGTCTCTAGGGCACTACATATC         588         ACT           2237244         GAAATAATGCTTCAGGGG         589         ACT           2237247         ATGCCTTCTAATGCATTCATTTTA         590         ACC           2237248         CCTAATCACATAAACCAGGAA         591         ACC           2237249         GAAAACAAGAGAGAGGGAAG         592         ACT           2239849         TGTGACTCCTCATGCTAATC         593         ACC           2239850         CCAGTCAATGCCAGAAA         594         ACT	2157545	GCTGGCATAAATGAAATTG	<u>578</u>	ACG
2157550         TTACATGATGAATATTATGGAAGT         581         ACT           2213838         TTCCAGCATGATTCTAAGACA         582         ACT           2213839         CAACTTGAGATACAGTAAAAATT         583         ACC           2213840         TGAAATGAATTCTCCAATAGAC         584         ACC           2227237         ACCCTCAAGTCCTTTTG         585         ACC           2237238         TCCCAGGACCCTAAAAAAGT         586         CGT           2237241         CAGTATTGCTAATGGGTGTTC         587         ACT           2237242         TGTCTCTAGGGCACTACATATC         588         ACT           2237244         GAAATAATGCTTCAGGGG         589         ACT           2237247         ATGCCTTCTAATGCATTCATTTTA         590         ACC           2237248         CCTAATCACATAAACCAGGAA         591         ACC           2237249         GAAAACAAGAGAGAGGGAAG         592         ACT           2239849         TGTGACTCCTCATGCTAATC         593         ACC           2239850         CCAGTCAATGCCAGAAA         594         ACT	2157546	GTGCCAGGTCCCACACT	<u>579</u>	ACT
2213838         TTCCAGCATGATTCTAAGACA         582         ACT           2213839         CAACTTGAGATACAGTAAAAATT         583         ACC           2213840         TGAAATGAATTCTCCAATAGAC         584         ACC           2227237         ACCCTCAAGTCCTTTTG         585         ACC           2237238         TCCCAGGACCCTAAAAAAGT         586         CG           2237241         CAGTATTGCTAATGGGTGTTC         587         ACT           2237242         TGTCTCTAGGGCACTACATATC         588         ACT           2237244         GAAATAATGCTTCAGGGG         589         ACT           2237247         ATGCCTTCTAATGCATTCATTTTA         590         ACC           2237248         CCTAATCACATAAACCAGGAA         591         ACC           2237249         GAAAACAAGAGAGGGAAG         592         ACT           2239849         TGTGACTCCTCATGCTAATC         593         ACC           2239850         CCAGTCAATGCCAGAAA         594         ACT	2157547	GTACACATACAATGATTTTACTC	<u>580</u>	ACT
2213839         CAACTTGAGATACAGTAAAAATT         583         ACC           2213840         TGAAATGAATTCTCCAATAGAC         584         ACC           2227237         ACCCTCAAGTCCTTTTG         585         ACC           2237238         TCCCAGGACCCTAAAAAAGT         586         CG           2237241         CAGTATTGCTAATGGGTGTTC         587         ACT           2237242         TGTCTCTAGGGCACTACATATC         588         ACT           2237244         GAAATAATGCTTCAGGGG         589         ACT           2237247         ATGCCTTCTAATGCATTCATTTTA         590         ACC           2237248         CCTAATCACATAAACCAGGAA         591         ACC           2237249         GAAAACAAGAGAGGGAAG         592         ACT           2239849         TGTGACTCCTCATGCTAATC         593         ACC           2239850         CCAGTCAATGCCAGAAA         594         ACT	2157550	TTACATGATGAATATTATGGAAGT	<u>581</u>	ACT
2213840         TGAAATGAATTCTCCAATAGAC         584         ACC           2227237         ACCCTCAAGTCCTTTTG         585         ACC           2237238         TCCCAGGACCCTAAAAAAGT         586         CGT           2237241         CAGTATTGCTAATGGGTGTTC         587         ACT           2237242         TGTCTCTAGGGCACTACATATC         588         ACT           2237244         GAAATAATGCTTCAGGGG         589         ACT           2237247         ATGCCTTCTAATGCATTCATTTTA         590         ACC           2237248         CCTAATCACATAAACCAGGAA         591         ACC           2237249         GAAAACAAGAGAGGGAAG         592         ACT           2239849         TGTGACTCCTCATGCTAATC         593         ACC           2239850         CCAGTCAATGCCAGAAA         594         ACT	2213838	TTCCAGCATGATTCTAAGACA	<u>582</u>	ACT.
2227237         ACCCTCAAGTCCTTTTG         585         ACC           2237238         TCCCAGGACCCTAAAAAAGT         586         CG           2237241         CAGTATTGCTAATGGGTGTTC         587         ACT           2237242         TGTCTCTAGGGCACTACATATC         588         ACT           2237244         GAAATAATGCTTCAGGGG         589         ACT           2237247         ATGCCTTCTAATGCATTCATTTTA         590         ACC           2237248         CCTAATCACATAAACCAGGAA         591         ACC           2237249         GAAAACAAGAGAGGGAAG         592         ACT           2239849         TGTGACTCCTCATGCTAATC         593         ACC           2239850         CCAGTCAATGCCAGAAA         594         ACT	2213839	CAACTTGAGATACAGTAAAAATT	<u>583</u>	ACG
2237238         TCCCAGGACCCTAAAAAAGT         586         CGT           2237241         CAGTATTGCTAATGGGTGTTC         587         ACT           2237242         TGTCTCTAGGGCACTACATATC         588         ACT           2237244         GAAATAATGCTTCAGGGG         589         ACT           2237247         ATGCCTTCTAATGCATTCATTTTA         590         ACC           2237248         CCTAATCACATAAACCAGGAA         591         ACC           2237249         GAAAACAAGAGAGGGAAG         592         ACT           2239849         TGTGACTCCTCATGCTAATC         593         ACC           2239850         CCAGTCAATGCCAGAAA         594         ACT	2213840	TGAAATGAATTCTCCAATAGAC	<u>584</u>	ACG
2237241         CAGTATTGCTAATGGGTGTTC         587         ACT           2237242         TGTCTCTAGGGCACTACATATC         588         ACT           2237244         GAAATAATGCTTCAGGGG         589         ACT           2237247         ATGCCTTCTAATGCATTCATTTTA         590         ACC           2237248         CCTAATCACATAAACCAGGAA         591         ACC           2237249         GAAAACAAGAGAGGGAAG         592         ACT           2239849         TGTGACTCCTCATGCTAATC         593         ACC           2239850         CCAGTCAATGCCAGAAA         594         ACT	2227237	ACCCTCAAGTCCTTTTG	<u>585</u>	ACG
2237242         TGTCTCTAGGGCACTACATATC         588         ACT           2237244         GAAATAATGCTTCAGGGG         589         ACT           2237247         ATGCCTTCTAATGCATTCATTTTA         590         ACC           2237248         CCTAATCACATAAACCAGGAA         591         ACC           2237249         GAAAACAAGAGAGGGAAG         592         ACT           2239849         TGTGACTCCTCATGCTAATC         593         ACC           2239850         CCAGTCAATGCCAGAAA         594         ACT	2237238	TCCCAGGACCCTAAAAAAGT	<u>586</u>	CGT
2237244         GAAATAATGCTTCAGGGG         589         ACT           2237247         ATGCCTTCTAATGCATTCATTITA         590         ACC           2237248         CCTAATCACATAAACCAGGAA         591         ACC           2237249         GAAAACAAGAGAGGGAAG         592         ACC           2239849         TGTGACTCCTCATGCTAATC         593         ACC           2239850         CCAGTCAATGCCAGAAA         594         ACC	2237241	CAGTATTGCTAATGGGTGTTC	<u>587</u>	ACT
2237247         ATGCCTTCTAATGCATTCATTTTA         590         ACC           2237248         CCTAATCACATAAACCAGGAA         591         ACC           2237249         GAAAACAAGAGAGGGAAG         592         ACC           2239849         TGTGACTCCTCATGCTAATC         593         ACC           2239850         CCAGTCAATGCCAGAAA         594         ACC	2237242	TGTCTCTAGGGCACTACATATC	<u>588</u>	ACT
2237248         CCTAATCACATAAACCAGGAA         591         ACC           2237249         GAAAACAAGAGAGGGAAG         592         ACT           2239849         TGTGACTCCTCATGCTAATC         593         ACC           2239850         CCAGTCAATGCCAGAAA         594         ACT	2237244	GAAATAATGCTTCAGGGG	<u>589</u>	ACT
2237249         GAAAACAAGAGAGGGAAG         592         ACT           2239849         TGTGACTCCTCATGCTAATC         593         ACC           2239850         CCAGTCAATGCCAGAAA         594         ACT	2237247	ATGCCTTCTAATGCATTCATTTTA	<u>590</u>	ACG
2239849         TGTGACTCCTCATGCTAATC         593         ACC           2239850         CCAGTCAATGCCAGAAA         594         ACT	2237248	CCTAATCACATAAACCAGGAA	<u>591</u>	ACG
2239850 CCAGTCAATGCCAGAAA 594 ACT	2237249	GAAAACAAGAGAGGGAAG	<u>592</u>	ACT
	2239849	TGTGACTCCTCATGCTAATC	<u>593</u>	ACG
2269646 CAGTTAGACTGAAACGCACA 595 ACT	2239850	CCAGTCAATGCCAGAAA	<u>594</u>	ACT
220000 00000000000000000000000000000000	2269646	CAGTTAGACTGAAACGCACA	<u>595</u>	ACT
2277084 TGTGTCATTTAAATCCTTCA 596 ACT	2277084	TGTGTCATTTAAATCCTTCA	<u>596</u>	ACT
2282853 GAGCTAGAAGAAACCTGAAAG <u>597</u> ACT	2282853	GAGCTAGAAGAAACCTGAAAG	<u>597</u>	ACT
2282854 GTTGGTGTCCAAATGGCA 598 ACT	2282854	GTTGGTGTCCAAATGGCA	598	ACT
		ACATGAAGACACTACCC	599	ACT
	2301513		<del> </del>	ACT
	2345808	GACATTTAGGTTATTTCCAAATTT		ACT
			602	ACT
				ACG

dbSNP rs#	Extend Primer	SEQ ID NO.	Term Mix
3752577	AGAAATAGATGGAGCCAAAAG	<u>604</u>	ACG
3777925	GGATGGGACTGAAACTC	<u>605</u>	ACT
3777926	CTCTGTAATTTTTCATGTATGATA	<u>606</u>	ACT
3777927	ATGAGCTCCCTTCACTC	<u>607</u>	ACG
3777928	TGAGCAAGCTAGAGAGTA	<u>608</u>	CGT
3777929	AAGTGGAGAGCATTTACAT	<u>609</u>	ACT
3777932	TTTGGCCGACTGAAATG	<u>610</u>	ACT
3777934	GGTCAGAGTGTCACTGGGCTACA	<u>611</u>	ACT
3777941	TCCCAAATTTCCTTTTCA	<u>612</u>	ACG
3777942	ACAAGTGAAGGTAGTATTGT	<u>613</u>	CGT
3798357	GCCTGGCATCTGCTAATC	<u>614</u>	ACT
3798359	GCAAAGGCAGAGACTAT	<u>615</u>	ACT
3822941	ACACACGATGTTTCTCCAG	<u>616</u>	ACG
3948760	ACAGTTTTATGAGACAGGTA	<u>617</u>	ACT

Please amend Table 19 on pages 102-104 as follows:

**TABLE 19** 

dbSNP rs#	Forward PCR primer	SEQ ID NO.	Reverse PCR primer	SEQ ID NO.
54144	ACGTTGGATGAGCAGCCATCACATGATCTG	618	ACGTTGGATGCCAATTGTGTCAAACATTTAATGAA	<u>619</u>
183535	ACGTTGGATGTGGTCTAAGTCCTGCACAAG	620	ACGTTGGATGGACAGAAAGAGTGTGCAGTC	<u>621</u>
236104	ACGTTGGATGCTCCCTGAACTTCCCATTTC	<u>622</u>	ACGTTGGATGCAGGGAGCTTACTACAAAGC	<u>623</u>
236105	ACGTTGGATGGGTAGCCACATTTGACACAG	<u>624</u>	ACGTTGGATGAAGTGGCCTGGAAACCAATG	625
236113	ACGTTGGATGGCTAATACATCCTAAAGGAAC	<u>626</u>	ACGTTGGATGCTTTATTGGAAATCTGTTCG	<u>627</u>
236119	ACGTTGGATGTTTCTTCTTGTCTCACAGGC	<u>628</u>	ACGTTGGATGCTGTAGATTTTCCTTTTGGC	<u>629</u>
236120	ACGTTGGATGCAGGATAATGATCATCCCTG	<u>630</u>	ACGTTGGATGACTGGCACAATTAGTGTCTG	<u>631</u>
236122	ACGTTGGATGTAGCTATCCCATTGTTGAGG	<u>632</u>	ACGTTGGATGGCTAAAGTCTGAAAACACTAG	<u>633</u>
236143	ACGTTGGATGCCTACCCCAAATAGGTAAAG	<u>634</u>	ACGTTGGATGAGGAAGTAAGTTTTGGGAGG	<u>635</u>
236145	ACGTTGGATGAGAATGTGAGCAAGGGATGC	<u>636</u>	ACGTTGGATGAATGGCACAGCTATCCTCAG	<u>637</u>
236146	ACGTTGGATGGTAAATCTGTATCTCCGCCC	<u>638</u>	ACGTTGGATGGACTTGATTACGTGACCTGG	<u>639</u>
236149	ACGTTGGATGCCAATGATCAAACTGAAATCG	<u>640</u>	ACGTTGGATGGGACTGGATTAGATGAATTC	<u>641</u>
236160	ACGTTGGATGCACCCACAGCTATCTGAGTT	<u>642</u>	ACGTTGGATGTCCAAAAGGAGTGGGTAGAG	<u>643</u>
236162	ACGTTGGATGAATGGAGTGTTTGCATGTTG	<u>644</u>	ACGTTGGATGCATGGATTTTTAGGACATGCG	<u>645</u>
236163	ACGTTGGATGGCCAAAAATAGCCTTTTCTC	<u>646</u>	ACGTTGGATGAAACAACATGCAAACACTCC	<u>647</u>
236164	ACGTTGGATGTTGCAAGCTGGTGTCACACA	<u>648</u>	ACGTTGGATGAGACCAGCCACTACTGATTC	<u>649</u>
236166	ACGTTGGATGCTCCTGTCATAGAATAGGCC	<u>650</u>	ACGTTGGATGGCATAGCACATGCTATTTGG	<u>651</u>
236167	ACGTTGGATGATCAAAGTCTTGTGCAGGAC	<u>652</u>	ACGTTGGATGGCACTTTAGGGACATTTGAC	<u>653</u>
236176	ACGTTGGATGGCATAGATAGCTTAATCATGG	<u>654</u>	ACGTTGGATGAAACCAATAGAAGCAGGTTG	<u>655</u>
236177	ACGTTGGATGCATTTGAGGATCGGAGTGAG	<u>656</u>	ACGTTGGATGCCCTGTGTCTGCAAATTTGG	657
236180	ACGTTGGATGCCTAGCACTTGGGAATTAGG	<u>658</u>	ACGTTGGATGGATGCGTGAAATAGATGCTC	<u>659</u>
236182	ACGTTGGATGCTCTCTAGTTCCTTTGTTGC	<u>660</u>	ACGTTGGATGATTTCAAAAGTGGTCTCCAC	661
236183	ACGTTGGATGTAAAAGAGAAATCCCACAGG	662	ACGTTGGATGCTCAGCAGATGTTAGTTTTC	663

dbSNP rs#	Forward PCR primer	SEQ ID NO.	Reverse PCR primer	SEQ ID NO.
236184	ACGTTGGATGCTAGGGACCCAGCAAATAAC	<u>664</u>	ACGTTGGATGACCATCTGAGGGAAATCCTG	<u>665</u>
379418	ACGTTGGATGTCTGGCCTCTAAGCTAAAGG	<u>666</u>	ACGTTGGATGCAGTGGTGTTGATAGATGGG	<u>667</u>
400735	ACGTTGGATGACCTTCTCAGGTTCACGTTC	<u>668</u>	ACGTTGGATGAAGGGCACCATGCCATTAAC	<u>669</u>
409035	ACGTTGGATGTGCTATGGGTATGCAAGGTG	<u>670</u>	ACGTTGGATGGGTTGTTGAAGGAGCGAGAG	<u>671</u>
440005	ACGTTGGATGATGGCTCTAAAAAGCTGTCC	<u>672</u>	ACGTTGGATGCAGCCTTCTTCCTGATACAG	<u>673</u>
446614	ACGTTGGATGTAAAGCCCAGGGCTAAAGAC	<u>674</u>	ACGTTGGATGAGAAGATGGCCAGAAAGGAG	<u>675</u>
451417	ACGTTGGATGCCAGAGTCATCGTTATCACC	<u>676</u>	ACGTTGGATGCCTCACTAAGGATTCAACCC	<u>677</u>
454422	ACGTTGGATGCAGCTTTTGAGGCACTTTCC	<u>678</u>	ACGTTGGATGAGCACCTTGCATACCCATAG	<u>679</u>
500277	ACGTTGGATGAGTTTCCCTACGTCTCTCTC	<u>680</u>	ACGTTGGATGAGTAACTCTAGCCTCTGCTC	<u>681</u>
540717	ACGTTGGATGCATCCAAAACCCAACAAATCC	682	ACGTTGGATGAGAGAGGTGTGTGACTTTTC	<u>683</u>
546106	ACGTTGGATGTTATAGCACTGATGGGCTCC	684	ACGTTGGATGCTGTGACATACTTTTCCAGG	<u>685</u>
571039	ACGTTGGATGATTCCTGTAGCAGGCAACTG	686	ACGTTGGATGGCTAGCTCTACTCTCTCTC	<u>687</u>
1039542	ACGTTGGATGTGAGGTTCTGTCTGAACACC	688	ACGTTGGATGTGGCTGCAATGGCTAACTTC	689
1039543	ACGTTGGATGATCTGACTCAGAAGAAGAGC	690	ACGTTGGATGGGCATTAATGGAGGTTATGC	<u>691</u>
1343180	ACGTTGGATGAGATGGCAACAGCAACACAG	692	ACGTTGGATGCCAACAGCAGCTTCACAATC	693
CHGB_SNP2	ACGTTGGATGAAATGGTATGTTTGTGTTCC	694	ACGTTGGATGTAATTTTTCCCCCCCAAATC	695
rs384578	ACGTTGGATGAAATGGTATGTTTGTGTTCC	696	ACGTTGGATGTAATTTTTCCCCCCCAAATC	697
rs742710	ACGTTGGATGAGAAAGTGAGGAAGAGAGGG	698	ACGTTGGATGATGAAATAGGCACGTGGCTC	699
rs742711	ACGTTGGATGATGAAATAGGCACGTGGCTC	700	ACGTTGGATGAGAAAGTGAGGAAGAGAGGG	701
rs881118	ACGTTGGATGTATAGCTGAAGCCTGCTTTC	702	ACGTTGGATGCAGTGAAGAGAAACACCTTG	703
rs910122	ACGTTGGATGAAGGTGTTTCTCTTCACTGC	704	ACGTTGGATGGGAGGGAGAACTATCAAA	705
CHGB_SNP1	ACGTTGGATGTCACTCTGAGGTCTTGGAGC	706	ACGTTGGATGTAAAGGGTTATCCAGGCGTC	707
180477	ACGTTGGATGGGAAGTAATTCTCTGGGCTG	708	ACGTTGGATGAAGTGATCCTCCCACCTCAG	709
236102	ACGTTGGATGCAGCCTGTTCTCTCTGAAAC	710	ACGTTGGATGGGATGCAAGAGGTTGTAGAG	711
236103	ACGTTGGATGCCTGTTTAAATCGTGGCTCC	712	ACGTTGGATGAAACATAAGGAAGCTGAGGC	713
236106	ACGTTGGATGCAAGCCTTTGCAGCTCTATC	714	ACGTTGGATGCCTCATAAGGGCCTTTGTAC	715
236107	ACGTTGGATGGAAGTTTACGTAAACTCTAG	716	ACGTTGGATGGTGTGTGGCTTATTGTAGAG	717
236108	ACGTTGGATGGTATTTACTGTTGAACCCAG	718	ACGTTGGATGATGTGGGTAAGTTGTGCACC	719
236109	ACGTTGGATGAGATTACAGGCACTAGCCAC	720	ACGTTGGATGTCTGGGCAACATGGTGAAAC	721
236110	ACGTTGGATGATCGATCCAATGTTGACTGC	722	ACGTTGGATGTTTCAGAACAAACCCCACAG	723
236111	ACGTTGGATGTTCAGGAAGCAGCAACCATC	724	ACGTTGGATGTATGCTGTGACCTCTCCAAC	725
236112	ACGTTGGATGAACGAGGTCAGGAGATCAAG	726	ACGTTGGATGCACGCCCGGCTAATTTTTTC	727
236114	ACGTTGGATGGAACCAAGGAAGTCTGACTC	728	ACGTTGGATGAAAGCTACCAGTCATGTGCC	729
236115	ACGTTGGATGATCAAAGTCCATACTGCAGG	730	ACGTTGGATGTATGATCGTAGGCACTGGAG	731
236116	ACGTTGGATGTGTTGTATTACCTGACCCTG	732	ACGTTGGATGAAGCAAACCACTGAGTGTCC	<u>733</u>
236117	ACGTTGGATGCAATGGTGTGATCTTGCCTC	734	ACGTTGGATGATTAGCCAAGTGTGGCAGTG	<u>735</u>
236118	ACGTTGGATGGGTTGAGTATCCCTAATCTG	736	ACGTTGGATGCTTTCAGTGTCGTGTCAGGG	737
236121	ACGTTGGATGCAAGCTATGTCACAGTTTAAG	738	ACGTTGGATGAGTCTTTGCCCTTAATGTGG	739
236123	ACGTTGGATGATAATAAATTTAGACTTCAC	740	ACGTTGGATGAAAATACTGGTGCGGCCAGA	<u>741</u>
236124	ACGTTGGATGGAATTTTGTTTGGCTCACGG	742	ACGTTGGATGATTGCTGCTGGAAGCTTACC	743
236125	ACGTTGGATGCCATGCCTGAGTTATTTGC	744	ACGTTGGATGATGGAGAAAGTAGATAGTAG	745
236148	ACGTTGGATGTAAGCCCAAGTGCTGTTGAG	746	ACGTTGGATGCTCAGAAGTCTGATGTGTATC	747
236151	ACGTTGGATGTTGGCCTTTAGACTCCTGGG	748	ACGTTGGATGAGAAGACACATAGCCGAGAG	749
236152	ACGTTGGATGAATAAAGGGTTATCCAGGCG	750	ACGTTGGATGTGGAGCCCTGTATTCTTCAC	<u>751</u>
236154	ACGTTGGATGTTCTGACAAGTTCCTGGCTG	752	ACGTTGGATGGCTGCATTAGTCAACCTACC	<u>753</u>
236155	ACGTTGGATGGGTAGGTTGACTAATGCAGC	754	ACGTTGGATGTGAGGTCCCGAACCAATTTC	<u>755</u>

dbSNP	Forward	SEQ ID	Q ID Reverse	
rs#	PCR primer	NO.	PCR primer	NO.
236158	ACGTTGGATGAAACTCCTGACCTCGTGATC	<u>756</u>	66 ACGTTGGATGCTCCTTAAGAAGATAGAGGC	
236159	ACGTTGGATGGTCTCAAACTCCTGACCTCG	<u>758</u>	ACGTTGGATGAAGAAGATAGAGGCAGCTGG	<u>759</u>
236161	ACGTTGGATGGATGTTGCCTCTAGGCTAGT	<u>760</u>	ACGTTGGATGCACCATCTGACCTGTGCTAC	<u>761</u>
236165	ACGTTGGATGAAAATTAGCCATGCGTGGTG	<u>762</u>	ACGTTGGATGTTCAAGCGGTTCTCCTGCCT	<u>763</u>
236168	ACGTTGGATGTCTATGTCTCCACTTGCATG	<u>764</u>	ACGTTGGATGACACATTTGCACACACACAC	<u>765</u>
236169	ACGTTGGATGGTGACTAGAATTTTTGTGTAC	<u>766</u>	ACGTTGGATGGTGTGCAAATGTGTATCC	<u>767</u>
236171	ACGTTGGATGAACCTCCCACTTTGGCTTTC	<u>768</u>	ACGTTGGATGGGTCCATTTAAAGCCTGGTG	<u>769</u>
236173	ACGTTGGATGATCAACCTGCACCACCAATC	<u>770</u>	ACGTTGGATGGCTAAGATGGAAGTTGAAGTG	<u>771</u>
236175	ACGTTGGATGTTCTCCATCACTGCATCAAG	<u>772</u>	ACGTTGGATGGTTATAGCCTGTATCGCAGC	<u>773</u>
236179	ACGTTGGATGCTAAATAACAGGTTTGACTC	<u>774</u>	ACGTTGGATGGAACATTGAGAGTATCTTAT	<u>775</u>
236181	ACGTTGGATGGTGAACATGTCTTTTCTGTAC	<u>776</u>	ACGTTGGATGGGTAGAACCACTGTTTTTCG	777
236185	ACGTTGGATGGGGTCACTTGAATTCAGGAG	<u>778</u>	ACGTTGGATGACTGCAACCACTGCCTCTTG	<u>779</u>
236187	ACGTTGGATGTAGTGAAACTCTGTCTCTGC	<u>780</u>	ACGTTGGATGACCTGCACCAACCTTTAACC	<u>781</u>
236189	ACGTTGGATGTGGATTTACAGAAAAACTGC	<u>782</u>	ACGTTGGATGCTGTGAGACACTAGGGATAC	<u>783</u>
364652	ACGTTGGATGTTTCTGCTGGGCTGTGATAG	<u>784</u>	ACGTTGGATGGGGAAATGCTCAGCATGTAC	<u>785</u>
394604	ACGTTGGATGTATTTTGGGATGGTGTGGGC	<u>786</u>	ACGTTGGATGGAACCAGGTCTTCCTTGATG	<u>787</u>
403727	ACGTTGGATGTCACTTGAACCCAGGAGATG	<u>788</u>	ACGTTGGATGGTTTTGAGACAGAGTTTCGC	<u>789</u>
446658	ACGTTGGATGTCACTGAGTTCAACTCCTTC	<u>790</u>	ACGTTGGATGGTTCCTGCTTTACCACTTCG	<u>791</u>
451571	ACGTTGGATGTTCTGGGTGGTTGCTCTCTG	<u>792</u>	ACGTTGGATGAAGTAATGGCACACTGGAGG	<u>793</u>
452749	ACGTTGGATGTCCTACTCCAGTATGACCTC	<u>794</u>	ACGTTGGATGGAAGTCCCAACCCCTAATAC	<u>795</u>
454328	ACGTTGGATGTGCAAACTGGTGCATCAGAG	<u>796</u>	ACGTTGGATGCCTGGTATTTTCATATCGCC	<u>797</u>
742710	ACGTTGGATGGCACATGGATATGGTGAAG	<u>798</u>	ACGTTGGATGTGCCTCTGTGATGGTGTCCC	799
742711	ACGTTGGATGGGCACATGGATATGGTGAAG	<u>800</u>	ACGTTGGATGAAATAGGCACGTGGCTCCCC	<u>801</u>
881118	ACGTTGGATGTATAGCTGAAGCCTGCTTTC	<u>802</u>	ACGTTGGATGTAGCAGTGAAGAGAAACACC	803
910122	ACGTTGGATGGTTTCTCTTCACTGCTATCT	<u>804</u>	ACGTTGGATGACACGCCATTCTGAGAAGAG	<u>805</u>
1005517	ACGTTGGATGTACTAATGTCAGTGGTAGAG	<u>806</u>	ACGTTGGATGTGAAGACACTGGCTGAAAAC	<u>807</u>
1394095	ACGTTGGATGTTCAGTGATCCAACTTCCGC	<u>808</u>	ACGTTGGATGCCAAACTCCTTGATTGGC	<u>809</u>
2206817	ACGTTGGATGGCAGAAACCCAGTGAAGTAG	<u>810</u>	ACGTTGGATGAAACCAGTTACTAACTGTAG	<u>811</u>
2268339	ACGTTGGATGATCCTGGAGATGTTATACCC	<u>812</u>	ACGTTGGATGCCTGGTGTTTAAGGCTCAAC	<u>813</u>
2300427	ACGTTGGATGAGATTACAGGCATGAGCCAC	<u>814</u>	ACGTTGGATGAAGTTAAATAAGCTCTTCTG	<u>815</u>
2326680	ACGTTGGATGAGGCTAATTCCTTCTCCTGG	<u>816</u>	ACGTTGGATGTCGTGCAACATCACTGTGTC	<u>817</u>
2423131	ACGTTGGATGATGCCTGCCTTACGAGAATG	<u>818</u>	ACGTTGGATGTGTCACTAGAATATGTGAAC	<u>819</u>
2876003	ACGTTGGATGGCAAAGACTAAGAGTCTGTAG	<u>820</u>	ACGTTGGATGCTGAGCCAGATTCTGACATT	<u>821</u>
3761873	ACGTTGGATGCTGTCCCTCTTAGAGCAATG	<u>822</u>	ACGTTGGATGCTATGAGCCTTTGACACAGC	<u>823</u>

Please amend Table 20 on pages 104-107 as follows:

TABLE 20

dbSNP rs#	Extend Primer	SEQ ID NO.	Term Mix
54144	CTGAAAGACACCATTTAT	<u>824</u>	CGT
183535	GTCCTGCACAAGACTTTGATA	<u>825</u>	ACG
236104	CCCATTTCATACCACCTATCA	<u>826</u>	ACG
236105	CTCCCTCCTCGAGACC	<u>827</u>	ACT

dbSNP	Extend	SEQ ID	Term
rs#	Primer	NO.	Mix
236113	GATCATTCATGAAACAGATTCTA	<u>828</u>	ACG
236119	TGTTCTCAAGGAAAAAAGAAAAA	<u>829</u>	ACT
236120	GATCATCCCTGGGAATGGTA	<u>830</u>	ACT
236122	GAGGCAGGGAATCAGCAATA	<u>831</u>	ACT
236143	ACCCCAAATAGGTAAAGATCTGT	<u>832</u>	ACT
236145	CTCCTGCACTGAGCTCCTAT	<u>833</u>	ACT
236146	TATCTCCGCCCTAAGAATACT	<u>834</u>	ACT
236149	GAAATATTAGAATTTAGAGGCAG	<u>835</u>	ACT
236160	GAGTTTTATGAGAAAGGGCAA	<u>836</u>	ACT
236162	GTTGTTTTAAAGTGTTGGTTGTAA	837	ACT
236163	CAATACATAGTGAAGCTTTGGG	<u>838</u>	ACT
236164	CTGGTGTCACACACACATGTA	839	ACT
236166	GGAACATCTCAGAAAAAAA	840	CGT
236167	CTTGTGCAGGACTTAGACCA	841	ACT
236176	ATAGGCTTTCTTGTGTATTTGCA	842	ACT
236177	AGTGAGGGGAAGCAGAGTC	843	CGT
236180	ACTTGGGAATTAGGTGGAGG	844	ACG
236182	GTTCAGAGATAATGCTGCTGATC	845	CGT
236183	GAAATCCCACAGGAACACAAT	846	ACT
236184	CCCAGCAAATAACAAGAATTGGCC	847	ACT
379418	CTTAAGCCAAGACAAACA	848	ACG
400735	TTCATCTTCCACCCTGGCC	849	ACT
409035	TGCTTTGCTTGCCTCCCACA	850	ACG
440005	GCTGTCCTTTTTACAAGGAAAT	<u>851</u>	ACT
446614	TAAAGACTGAAGCTTTCACAGT	852	ACT
451417	CGTTATCACCATTGGGCTTTA	<u>853</u>	CGT
. 454422	GATCCTTCTCACTTACTGTTC	<u>854</u>	ACT
500277	GATTATGCCCTGAGGTCTTTTG	<u>855</u>	ACT
540717	AACCCAACAAATCCTAGGGC	<u>856</u>	ACG
546106	GATGGGCTCCCCATATGAC	<u>857</u>	ACT
571039	TGTAGCAGGCAACTGAGCAGGAGA	<u>858</u>	CGT
1039542	GAACACCCTCCAGCACAAG	<u>859</u>	ACT
1039543	AGAAGAGCTTTCATCTGTGTG	<u>860</u>	ACT
1343180	CACAGCCCTCCATTACAGC	<u>861</u>	ACT
CHGB_SNP2	GTATGTTTGTGTTCCATTTGCA	<u>862</u>	ACT
rs384578	GTATGTTTGTGTTCCATTTGCA	<u>863</u>	ACT
rs742710	GAAGAGAGGGCCTTGAGC	<u>864</u>	ACG
rs742711	TCCCCTGCCTCTGTGATGG	<u>865</u>	ACG
rs881118	CTGAAGCCTGCTTTCTTCAT	<u>866</u>	ACT
rs910122	CTTCACTGCTATCTTCCCCT	<u>867</u>	ACG
CHGB_SNP1	TCTTGGAGCCCTGTATTC	<u>868</u>	ACG
180477	GTGGCTCACGCCTATAA	<u>869</u>	CGT
236102	GTTCTCTGAAACCTGTTA	<u>870</u>	ACT
236103	CATGCACCAGCTGTGTG	<u>871</u>	ACT

dbSNP	Extend	SEQ ID	Term Mix
rs#	Primer	NO.	
236106	GAACATTCCAGGCAAAC	872	ACT
236107	GTTCTGGTAAAAAAAAAGTTTG	<u>873</u>	CGT
236108	CTGTTGAACCCAGAAATATC	<u>874</u>	ACT
236109	CACTAGCCACCACGCCC	<u>875</u>	ACT
236110	CAATGTTGACTGCATTGACT	<u>876</u>	ACT
236111	GTTCTGAGGTTACCAGA	<u>877</u>	ACT
236112	ACCATCCTAGCTAACACG	<u>878</u>	ACT
236114	AATCACAAGTACCTCGAATAC	<u>879</u>	ACT
236115	AGGTAAGTGGCAGAACT	<u>880</u>	ACT
236116	TCAGGCAAGCACAGTACAAA	<u>881</u>	ACG
236117	GCCTCCCAAGTAGCTGG	<u>882</u>	ACT
236118	CCCTAATCTGAAAATCTGAAATCT	<u>883</u>	ACT
236121	AAGAATTTTCTTATTCAACTGTC	<u>884</u>	ACT
236123	CTTCACTAAATAAAAATGTGTCC	<u>885</u>	ACG
236124	GTTTGGCTCACGGAATTAT	<u>886</u>	ACG
236125	TTTAACTCCTAGCTTTTAAAGA	887	ACG
236148	AATGTGGCTGGTCCGATCTG	888	ACT
236151	ATTCTCCTGGCTCCCTG	889	ACG
236152	TTATCCAGGCGTCCAGG	890	ACT
236154	TGATGCCACTGGTCAGG	891	ACT
236155	AATTCCCCTTTGCACTCAT	892	ACT
236158	TTACAACTGTAAGCCACCGC	893	ACT
236159	CCTGACCTCGTGATCTG	894	ACG
236161	CTCTAGGCTAGTATTAATTTTTGT	895	ACT
236165	TAACGCCTGTAATCCCA	896	ACT
236168	ACTTGCATGTGTATGTATATCT	897	ACT
236169	ATGTCTTTTCCCCCTCT	898	ACT
236171	AAGTGCTGGGATTACAGATA	899	ACG
236173	TTGCTCCCTCTCCCCTT	900	ACT
236175	CACTGCATCAAGATGGGCC	901	ACG
236179	CAGGTTTGACTCAAAACTTTAA	902	ACG
236181	ATGTCTTTTCTGTACTGGATA	903	ACT
236185	TACTGAGGAGGCTGAGG	904	ACG
236187	AACTCTGTCTCTGCAAAAAA	905	ACT
236189	CAGAAAACTGCACAAAA	906	ACT
364652	CTGTGATAGGAAAAAGGAA	907	ACT
394604	CCAGCAGAGGCAAAAATAAGA	908	ACG
403727	TGCCACTGCACTCCAGCCT	909	ACT
446658	AGGAAAAGAGAGGCAAAC	910	ACT
451571	GCTGTCTTCATTCTCTTGT	911	ACG
452749	CCTATTTTCAAGTCAGGT	912	ACG
454328	CCTAAACAGCAGTTTTAGTACAT	913	ACG
742710	AGAGAGGGCCTTGAGC	914	ACG
742710	TGAGCCGGGAAAGGGAC	915	ACT
172111	16	1 212	Seria

dbSNP rs#	Extend Primer	SEQ ID NO.	Term Mix
881118	TGAAGCCTGCTTTCTTCAT	<u>916</u>	ACT
910122	CTTCACTGCTATCTTCCCCT	917	ACG
1005517	GGTAGAGAATGTAATAACAGT	<u>918</u>	ACT
1394095	ACGAGAGGGGGGGGGG	<u>919</u>	ACT
2206817	TTAGAGCAGGGCAGGGG	920	ACT
2268339	CAGAATGCTGAGATGGC	<u>921</u>	ACT
2300427	CACCCGGCCGGGAAAAT	922	ACT
2326680	TGGAATTTGAGAAGGCCTG	923	ACT
2423131	GCCTTACGAGAATGTTATTT	924	CGT
2876003	AGAGTCTGTAGTCCCAA	<u>925</u>	ACT
3761873	TGTATTTTCCATAGTAATTTGCTC	<u>926</u>	ACT

Please amend Table 22 on page 110 as follows:

TABLE 22

dbSNP	First	SEQ ID	Second	SEQ
rs#	PCR primer	NO.	PCR primer	ID NO.
742710	ACGTTGGATGGACATGAAATAGGCACGTGG	<u>927</u>	ACGTTGGATGAGAAAGTGAGGAAGAGAGGG	<u>928</u>
236110	ACGTTGGATGTCACTCTGTTTCTACTAACC	<u>929</u>	ACGTTGGATGATCGATCCAATGTTGACTGC	930

Please amend Table 23 on page 110 as follows:

**TABLE 23** 

dbSNP rs#	Extend Primer	SEQ ID NO.	Term Mix
742710	AGAGAGGGCCTTGAGC	<u>931</u>	ACG
236110	AATGTTGACTGCATTGACT	<u>932</u>	ACT

Please amend Table 26 on pages 112-113 as follows:

**TABLE 26** 

dbSNP rs#	Forward PCR primer	SEQ ID NO.	Reverse PCR primer	SEQ ID NO.
752373	ACGTTGGATGTATCACAAGAACAGCATGGG	<u>933</u>	ACGTTGGATGATGGTTTCTGTAATCCCCCC	<u>934</u>
763470	ACGTTGGATGAAGAGGAGTGGCTGATAATG	<u>935</u>	ACGTTGGATGAAGCAGAAAACTTTGTGCCG	<u>936</u>
763471	ACGTTGGATGGGCAGGTCATGGATTTATTG	<u>937</u>	ACGTTGGATGCATCATTCCTCTGTGAGGCG	<u>938</u>
763471	ACGTTGGATGGTGAAGAGCTCTGAAATGCC	<u>939</u>	ACGTTGGATGTAACTCCTGTGTGGCTTTCT	940
899011	ACGTTGGATGAAGGTGGAGCCTGCCTCAAG	941	ACGTTGGATGAGCTTTGCACCCTGTGATGC	942

dbSNP	Forward	SEQ ID	Reverse	SEQ ID
rs#	PCR primer	<u>NO.</u>	PCR primer	NO.
899011	ACGTTGGATGAAGGTGGAGCCTGCCTCAAG	<u>943</u>	ACGTTGGATGAGCTTTGCACCCTGTGATGC	944
922359	ACGTTGGATGTCAAGCGATCCTCTTCAGCC	<u>945</u>	ACGTTGGATGATTCATTCCAAGACCGGGTG	<u>946</u>
930672	ACGTTGGATGGTGGGTTACTTGGTCCATAC	<u>947</u>	ACGTTGGATGACAGAGCAAGACCTTCTCTC	<u>948</u>
936513	ACGTTGGATGGTATGAAGTTCTTTGCAGAGT	<u>949</u>	ACGTTGGATGTACTACTGCACTCCAGCCTG	<u>950</u>
1375999	ACGTTGGATGCCATTCTTTTACCTTGAACC	<u>951</u>	ACGTTGGATGCAGAGACTTGCAGAATGGAC	<u>952</u>
1376000	ACGTTGGATGATAGCTGATGGTGTGCTGAG	<u>953</u>	ACGTTGGATGAAGCTTGCCTCCCAAGTTAG	<u>954</u>
1376001	ACGTTGGATGCAAACAATCCCATTACACAG	<u>955</u>	ACGTTGGATGCAGTACAACAGGGTGGCTATC	<u>956</u>
1450270	ACGTTGGATGCCATATCACATGGATATGAGG	<u>957</u>	ACGTTGGATGCATGGCTTCTCTTACACCTG	<u>958</u>
1450273	ACGTTGGATGGCTGCATATAAGAGACACATG	<u>959</u>	ACGTTGGATGGCCACTCCAGCTTTCTTTTG	<u>960</u>
1450274	ACGTTGGATGTGAGAGGAAGCCTGGTGTTG	<u>961</u>	ACGTTGGATGAAGCTTGCCTCCCAAGTTAG	<u>962</u>
1562781	ACGTTGGATGTATGTCTCCTGCCTTCTTCC	<u>963</u>	ACGTTGGATGGGAAAGAAGCTTGATGTGGC	<u>964</u>
2071019	ACGTTGGATGGGTAAACAACTGACCCATCC	<u>965</u>	ACGTTGGATGCCTGGGAAATAACCATGAGC	<u>966</u>
2071020	ACGTTGGATGAATTCACAGCTAAGCCTCCC	<u>967</u>	ACGTTGGATGTTCAGCTCCAGCTGCATGTT	<u>968</u>
2198008	ACGTTGGATGGTAGAAGTTTAGTATATGATG	<u>969</u>	ACGTTGGATGCCCTGTCATTTCAAATACCG	<u>970</u>
2198009	ACGTTGGATGCTTGTGCCAATCCCACAATG	<u>971</u>	ACGTTGGATGGCAGAAGTCTAGCCAAGAAC	<u>972</u>
2198010	ACGTTGGATGCTTGTGCCAATCCCACAATG	<u>973</u>	ACGTTGGATGAATGCAGAAGTCTAGCCAAG	<u>974</u>
2403330	ACGTTGGATGTAACTCTGAGACCCAAGGAC	<u>975</u>	ACGTTGGATGCCAGACAGTTGTGTGTTGAC	<u>976</u>
2923115	ACGTTGGATGGGATTACCCTAAGGATCCAC	<u>977</u>	ACGTTGGATGAGAGGAATTCAGTTGCTGCC	<u>978</u>
2923117	ACGTTGGATGTTGAGTCCAAGAGGTTGAGG	<u>979</u>	ACGTTGGATGAGACAGTCTTGCTCTGTCAC	<u>980</u>
2957666	ACGTTGGATGTACTTGGGAGACTGAGGTAG	<u>981</u>	ACGTTGGATGGCATAGTGGTGTGATCATGG	<u>982</u>
2957667	ACGTTGGATGAGAATGGTCTTTCCCACTCC	<u>983</u>	ACGTTGGATGATGGATTACGGAAGGAATAC	<u>984</u>
2957669	ACGTTGGATGTACTGAGACTCCCAGCATTG	<u>985</u>	ACGTTGGATGGTGTGCAGCTTAGTAAGTGC	<u>986</u>
2957670	ACGTTGGATGTCATGTGATTCTCCTGCCTC	<u>987</u>	ACGTTGGATGGTGAAACCCCGTCTCTACTA	<u>988</u>
2957675	ACGTTGGATGAGAATGACTTGGGTTTTGGG	<u>989</u>	ACGTTGGATGCAGTGAGTTGTGACAGCACC	990
2957677	ACGTTGGATGGTCTTTCTCAATCCCAGCAC	<u>991</u>	ACGTTGGATGACGAGATCTCCTTGTGTTGC	992
2957678	ACGTTGGATGAAGACCTCAGGATGTGATGC	<u>993</u>	ACGTTGGATGATGACCCCGTTTCTTTGCAC	<u>994</u>
2957679	ACGTTGGATGAGTTCGTCAGAGAGATGTCC	<u>995</u>	ACGTTGGATGGAGCACATGGATTCACAGAG	996
3741043	ACGTTGGATGGACATCAGAAGCTAATTGGG	<u>997</u>	ACGTTGGATGCTTCTTAATGGTAGGGCCAG	998
3741044	ACGTTGGATGTTTGTTATGCAGAGGTGGCC	<u>999</u>	ACGTTGGATGTAGATGGGCTCTTCTTGGAC	1000
3741045	ACGTTGGATGAACTGAGCTTCAGACTTCCC	<u>1001</u>	ACGTTGGATGTCAGACCTGTAGATGGGCTC	<u>1002</u>
3891547	ACGTTGGATGGCCATCAAGTTTGTGGCAAT	<u>1003</u>	ACGTTGGATGAAGCTATATGGAGCCCAAGG	1004

Please amend Table 27 on pages 113-114 as follows:

**TABLE 27** 

dbSNP rs#	Extend Primer	SEQ ID NO.	Term Mix
752373	GGGACCAGGTGGAGATAA	<u>1005</u>	ACG
763470	AGAAAACTTTGTGCCGTTTTCT	<u>1006</u>	ACT
763471	CCAGGCAGCAACTCCCT	<u>1007</u>	ACT
763471	CTCCAAGCAGTAAAGATGTTC	<u>1008</u>	CGT
899011	TTGGTTTTAGAGGATTGCTCC	<u>1009</u>	ACG
899011	GGTTTTAGAGGATTGCTCC	<u>1010</u>	ACG
922359	TCATGCCTATAATCCAAGCA	<u>1011</u>	ACT

dbSNP rs#	Extend Primer	SEQ ID NO.	Term Mix
930672	AAAAGCAAGAACAACAGCA	1012	CGT
936513	AGACAGGGTGAGACCTC	<u>1013</u>	ACT
1375999	TATGCTGCATATAAGAGACACAT	1014	ACT
1376000	ACATATTTCTGGTCTCCA	<u>1015</u>	ACT
1376001	ACAGGGTGGCTATCATTAAC	<u>1016</u>	ACT
1450270	TGTGAACTGAAAAGTCAAG	<u>1017</u>	ACT
1450273	CTTGAACCTATTTCTGTTTTT	<u>1018</u>	ACT
1450274	CTCCCAAGTTAGATTGGTTA	<u>1019</u>	ACT
1562781	CTTGATGTGGCTGAAGT	<u>1020</u>	CGT
2071019	GTGCTGTTGAAATCCTGGG	<u>1021</u>	ACT
2071020	AACCCTTTGTCAGCTGAA	<u>1022</u>	ACT
2198008	CAGGCTTTTGGCTAAGATCAAG	<u>1023</u>	ACT
2198009	AGTGAAGAATTTTCCCTATTAGAT	<u>1024</u>	ACT
2198010	AAGTCTAGCCAAGAACATTT	<u>1025</u>	ACT
2403330	CTCCCACTCCTCTCATCAG	<u>1026</u>	ACT
2923115	GTTGCTGCCCGCTTTCC	<u>1027</u>	ACG
2923117	CTCTGTCACCCATGCTGGA	<u>1028</u>	ACT
2957666	ACCTCCTGGGCTCAAGC	<u>1029</u>	ACT
2957667	GGAAGGAATACTAAAGAACAA	<u>1030</u>	CGT
2957669	AGTAAGTGCTGTGATGCACC	<u>1031</u>	ACT
2957670	TAGCTGAGCATGGTGGC	<u>1032</u>	ACT
2957675	AGCATGGGTGACAGAGC	<u>1033</u>	ACT
2957677	TGTGTTGCCCAGACTAG	<u>1034</u>	ACT
2957678	ACTCCCTGGCCTCCCCT	<u>1035</u>	ACG
2957679	TCACAGAGCTGCCAGGG	<u>1036</u>	ACT
3741043	CAAAATTCTCCTGCCAC	<u>1037</u>	ACT
3741044	GGGGAAAGGGAAGTCTG	<u>1038</u>	CGT
3741045	TAGATGGGCTCTTCTTG	<u>1039</u>	ACT
3891547	TATGGAGCCCAAGGATGACC	<u>1040</u>	ACT

Please amend Table 30 on pages 118-119 as follows:

TABLE 30

dbSNP rs#	Forward PCR primer	SEQ ID NO.	Reverse PCR primer	SEQ ID NO.
726215	ACGTTGGATGAACTGAGCCCCATGAAATGC	1041	ACGTTGGATGAAAACAGCAATTGAGAACAC	1042
966783	ACGTTGGATGCTCCTGAATTTTAGCCATAC	<u>1043</u>	ACGTTGGATGTACGCAATAGTTCCTGGGAG	1044
998329	ACGTTGGATGGAAGAGCACATTATTTGCTGG	<u>1045</u>	ACGTTGGATGACACACTGGTGTTTTGTCAG	1046
1387472	ACGTTGGATGGAAAGGCCTTGAATTGGAAC	<u>1047</u>	ACGTTGGATGGTTCTGCTAGTGTCATCTTC	<u>1048</u>
1484119	ACGTTGGATGGCAGCTACAATCATAAAGGG	<u>1049</u>	ACGTTGGATGTGTGCCCTTAATAATGGTTG	1050
1484120	ACGTTGGATGATGGTCATGGCATCCAGTTC	<u>1051</u>	ACGTTGGATGGGCTGGTTTCTGACACTATC	1052
1489479	ACGTTGGATGGGAGCATCAGTCATTTTGGG	<u>1053</u>	ACGTTGGATGCACCAGGACATAACATGACG	1054

dbSNP	Forward	SEQ ID	Reverse	SEQ ID
rs#	PCR primer	NO.	PCR primer	NO.
1489480	ACGTTGGATGGGGTTGTGGAGAATCATTAC	<u>1055</u>	ACGTTGGATGGGTGGCAGTAATCTTCACTT	<u>1056</u>
1489481	ACGTTGGATGTCTCTGCAGTTGAGGAGATG	<u>1057</u>	ACGTTGGATGTTGGGAAAGGCCATCAAGTC	<u>1058</u>
1489482	ACGTTGGATGCTCTGGATAAAAGACTCAGC	<u>1059</u>	ACGTTGGATGCCCTTCCAACAGCTATCTGG	<u>1060</u>
1489483	ACGTTGGATGTTGGTTTGCTATCAATGAAG	<u>1061</u>	ACGTTGGATGGATAGGTGTACACATATAGC	<u>1062</u>
1489486	ACGTTGGATGAAAAAACACACCACAGCCCC	<u>1063</u>	ACGTTGGATGCTTCGTATTTGGCTCTGACC	<u>1064</u>
1552280	ACGTTGGATGATGAAAAGTGACACCCATCC	<u>1065</u>	ACGTTGGATGTCTGAAGCTGTTGAATCAGG	<u>1066</u>
1565288	ACGTTGGATGTAGCCAATTGGTGAACACTC	<u>1067</u>	ACGTTGGATGCTGCCAGTCATAAGGCAAAG	<u>1068</u>
1844334	ACGTTGGATGGCCAAGGAAACTAATTCCTG	<u>1069</u>	ACGTTGGATGCACTTTGGAAGACAGTTCGG	<u>1070</u>
1872203	ACGTTGGATGGTTGCATTAGCTGTTATTCTC	<u>1071</u>	ACGTTGGATGCCAGCAATTCTATTTCAGAG	<u>1072</u>
1905520	ACGTTGGATGCATGGTTTATACTTACTTACG	<u>1073</u>	ACGTTGGATGGTTTATTCCTGTTTCCACAC	1074
2029395	ACGTTGGATGGGAGGGAGACAAAGATTCAC	<u>1075</u>	ACGTTGGATGGCAACAGTTTCACCTTTGGC	<u>1076</u>
2029397	ACGTTGGATGCTCACAGTCCTGAAGACTTG	<u>1077</u>	ACGTTGGATGTGGAAGTGAAGGAGAGAGC	<u>1078</u>
2046777	ACGTTGGATGGGACTTCAAATATGGTTCAC	<u>1079</u>	ACGTTGGATGTTAAGCCTGGGACTTTTGGG	<u>1080</u>
2046778	ACGTTGGATGGTTCCCTTCCCCCATAAAAC	<u>1081</u>	ACGTTGGATGCATGAAGCCTTATGCTTGAG	<u>1082</u>
2054708	ACGTTGGATGCTAGGCATATCATGCCTCTG	<u>1083</u>	ACGTTGGATGTTGAGCTCACTGTTACCTGC	<u>1084</u>
2078403	ACGTTGGATGTGTGCTCAGGATCGACAGAC	<u>1085</u>	ACGTTGGATGACTCGAGACAACCTACAAGG	<u>1086</u>
2086832	ACGTTGGATGCTTTTGAGCATCACATTCCTC	<u>1087</u>	ACGTTGGATGTGCCTAAGCACTGTATAACC	<u>1088</u>
2129108	ACGTTGGATGAACTCCCAGTAAGTCCTTCC	<u>1089</u>	ACGTTGGATGACTCAGGCAGTAACTCCAAC	<u>1090</u>
2129111	ACGTTGGATGTACACTTTTCCCGCAAGACC	<u>1091</u>	ACGTTGGATGGTCATGGACATCTACAGTATC	<u>1092</u>
2170850	ACGTTGGATGGAAGGCCAATGCAAGGATAC	<u>1093</u>	ACGTTGGATGAAGAACACACAAAAAAAAT	<u>1094</u>
2279472	ACGTTGGATGGAGAAGAGCATTGGTTGCTG	<u>1095</u>	ACGTTGGATGTGCCCACAAGTGCTATCTAC	<u>1096</u>
2291304	ACGTTGGATGGTCTCAGGAAGGTTTAGAGG	<u>1097</u>	ACGTTGGATGAAAAGACAAACGATATGGCC	<u>1098</u>
2291305	ACGTTGGATGCATGATTTCAAAATCATGTTC	1099	ACGTTGGATGGAGATGTACAGTATGAGTCC	<u>1100</u>
2291306	ACGTTGGATGCAGCGACTAGTCATTAACCG	<u>1101</u>	ACGTTGGATGCAGTTGGTTTCAACTCTGCC	1102
2291309	ACGTTGGATGCATTGTTGTTCCTACCATTC	<u>1103</u>	ACGTTGGATGAAAGTGGTAAAGGAGAGGCG	1104
2291310	ACGTTGGATGGTGCTTGATACTTGGCCTAC	<u>1105</u>	ACGTTGGATGCAACTGGAAATTGCCGAAGC	<u>1106</u>
2291311	ACGTTGGATGTCAACATTTACTCCTAGCTC	<u>1107</u>	ACGTTGGATGATTTTGGGCTGTGGTCTTCC	<u>1108</u>
2291312	ACGTTGGATGTGTATTCTCCTGCATCGCTC	<u>1109</u>	ACGTTGGATGTCCAAGTTCAAGAACGACAC	<u>1110</u>
2291313	ACGTTGGATGTTCGAGTTTACCGTATGGTG	<u>1111</u>	ACGTTGGATGGATCACAGACAGGTCAGTTG	1112
2306636	ACGTTGGATGCTGAGACCAGTCTGTGTTTG	<u>1113</u>	ACGTTGGATGGTTTCCCATGACACTGTTCC	<u>1114</u>
2306637	ACGTTGGATGCTACTACTATTTCTGGAGTC	<u>1115</u>	ACGTTGGATGCTTATGCATTTCAACTGCCAC	<u>1116</u>
2366911	ACGTTGGATGGTAGATGCTTGAATCAATAAAG	<u>1117</u>	ACGTTGGATGATAGCAGCTCCAGAACTAGG	<u>1118</u>
2366912	ACGTTGGATGGAACTGTTGTTGAATGGGAC	<u>1119</u>	ACGTTGGATGCAATACTTGTAAAATAGCAGC	<u>1120</u>
2366913	ACGTTGGATGCTATCTGTATTCTCATGGCTG	<u>1121</u>	ACGTTGGATGTTACCTAGTTCTGGAGCTGC	<u>1122</u>
3769858	ACGTTGGATGCTACATGTCCATGGTTTGATG	1123	ACGTTGGATGGCATCAACCTTTATGCCAAG	<u>1124</u>
3769860	ACGTTGGATGGTATACAGAATATTGCATGCC	<u>1125</u>	ACGTTGGATGGAACATCATTGAAGGTAAAG	<u>1126</u>
3769863	ACGTTGGATGCAAGGATTTATTACATGCTG	<u>1127</u>	ACGTTGGATGGTCATCAGGAGAAAGTAAGC	<u>1128</u>
3816782	ACGTTGGATGGAGGAAACCAGAGCTTCAAG	<u>1129</u>	ACGTTGGATGCAGCACGCTGTTTCTCAATG	<u>1130</u>
3816849	ACGTTGGATGAACCAGCTCACCTCAGGAAC	<u>1131</u>	ACGTTGGATGTTTGTGGTGCCCATTCAAAC	<u>1132</u>

Please amend Table 31 on pages 119-120 as follows:

**TABLE 31** 

dbSNP	SEQ ID	Extend	Term
rs#	NO.	Primer	Mix
726215	<u>1133</u>	TTGAGAACACAGGATGC	ACT
966783	<u>1134</u>	CTCCCATTTTGGTCTTG	ACG
998329	<u>1135</u>	GGTGTTTTGTCAGTACAATT	ACG
1387472	<u>1136</u>	ACTACAAACTCTTCCTTACC	CGT
1484119	<u>1137</u>	GTTGTTTATGTTATGTTATGTGTT	ACT
1484120	<u>1138</u>	TGTGCCTCAGTTTCTCC	CGT
1489479	<u>1139</u>	GACAGCTGTAATTGTAGACC	ACT
1489480	<u>1140</u>	CTCAATCACATTTACCCTC	ACT
1489481	<u>1141</u>	TCTGATTGTTCCATTAATATCTG	ACT
1489482	<u>1142</u>	CAGCTATCTGGAAATCTTGTTTGA	CGT
1489483	1143	GTGTACACATATAGCAACCTCA	ACT
1489486	1144	CTCTGACCTGTGAGCTAC	ACT
1552280	<u>1145</u>	GCTGTTGAATCAGGATTTGATT	ACG
1565288	1146	GGCAAAGAAACACTAGAAA	ACG
1844334	1147	CAGTTCGGCAGTTTCTT	ACT
1872203	1148	AAAAATCATGAAAAGGAGCATG	CGT
1905520	1149	ACAAGTCTTTTCATGGTC	ACG
2029395	1150	CAAAATGAAGGAACACTTATCA	ACG
2029397	1151	AGCTCTGTTGGCACTTT	ACT
2046777	1152	GAGCCTGATTATTTGTTTGGGTA	ACG
2046778	1153	CTGTCATGATTGACAGGTCC	ACT
2054708	1154	CCTGGGCCTGGAAGGCAAC	ACG
2078403	1155	GGCTGGAGCAAGAATTA	ACG
2086832	1156	CAATGTAATCCTTGGATAGAT	CGT
2129108	1157	CAACTACATAGTCAGACTTT	ACT
2129111	1158	TATACGCAATAGTTCCTGGG	ACT
2170850	1159	GAACACACAAAAAAATTTAATCA	ACT
2279472	1160	CTCTTTAAACCTGCATTTTC	ACT
2291304	1161	CGATATGGCCATTTTGG	ACT
2291305	1162	CATATTCACACAATGGGAAAA	CGT
2291306	1163	CTGCCAACTATCAGCTT	ACT
2291309	1164	GCGAGACCATGGCATATAACA	ACT
2291310	1165	AACTTACACGTTTGTTGCTA	ACT
2291311	1166	GTGGTCTTCCGGATATCA	ACG
2291312	1167	CGACACAAATATGTAGTGGA	ACT
2291313	1168	TGTCTTGCTACATTCCAGT	ACT
2306636	1169	TCCAGTAAAATGGTTCCATAAGA	ACT
2306637	1170	TCAACTGCCACAAAATG	ACT
2366911	1171	TTCCTTTGTCCCATTCA	ACT

dbSNP rs#	SEQ ID NO.	Extend Primer	Term Mix
2366912	1172	TGTAAAATAGCAGCTCCAGAA	CGT
2366913	<u>1173</u>	ATTCTAAATGGAAAAAGAGCCA	ACG
3769858	<u>1174</u>	TGCCCTGAATGTGCCTC	ACT
3769860	<u>1175</u>	GGATAAGCATATGTAACTTTACG	CGT
3769863	<u>1176</u>	AAGTAAAAAGGACATAAAAACCT	ACT
3816782	<u>1177</u> -	GTTGATGGAACAACATAAAA	CGT
3816849	1178	GCCCATTCAAACATAAAG	ACT